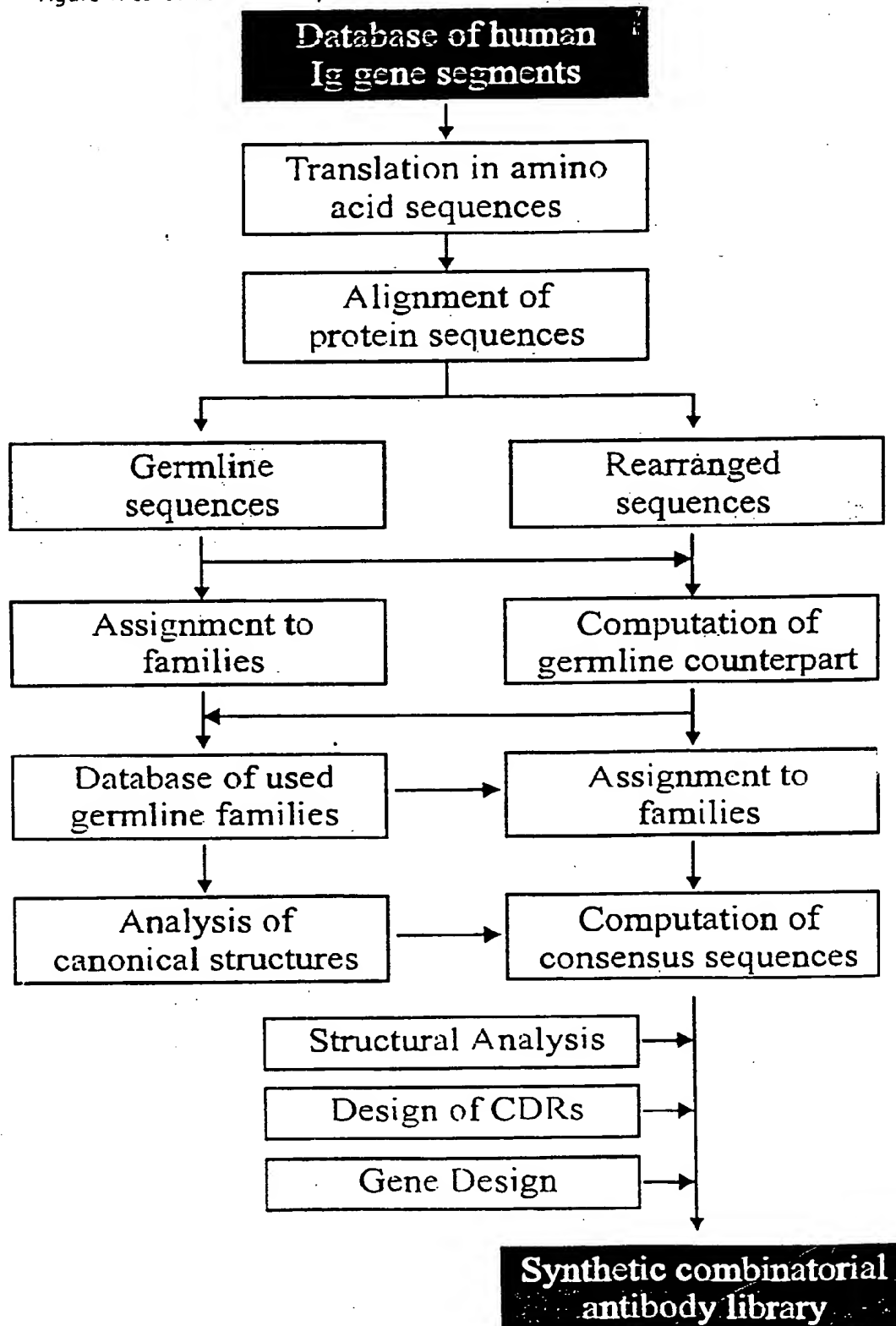


Figure 1: construction of a synthetic human antibody library based on consensus sequences



framework 1		CDRI	
	1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28		
Vk1	D I Q M T Q S P S S L S A S V G D R V T I T C R A S Q - -		
Vk2	D I V M T Q S P L S L P V T P G E P A S I S C R S S Q S L L		
Vk3	D I V L T Q S P A T L S L S P G E R A T L S C R A S Q S - -		
Vk4	D I V M T Q S P D S L A V S L G E R A T I N C R S S Q S V L		

CDRI			framework 2										CDR II																
D	F	F	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54
-	-	-	G	I	S	S	Y	L	A	W	Y	Q	Q	K	P	G	K	A	P	K	L	L	I	Y	A	A	S	S	L
H	S	-	N	G	Y	N	Y	L	D	W	Y	L	Q	K	P	G	Q	S	P	Q	L	L	I	Y	L	G	S	N	R
-	-	-	V	S	S	S	Y	L	A	W	Y	Q	Q	K	P	G	Q	A	P	R	L	L	I	Y	G	A	S	S	R
Y	S	S	N	N	K	N	Y	L	A	W	Y	Q	Q	K	P	G	Q	P	P	K	L	L	I	Y	W	A	S	T	R

Figure 2A: VL kappa consensus sequences

CDRII		framework 3																																																										
Vk1	55	Q	56	S	57	G	58	V	59	P	60	S	61	R	62	F	63	S	64	G	65	S	66	G	67	S	68	G	69	T	70	D	71	F	72	T	73	L	74	T	75	I	76	S	77	S	78	L	79	Q	80	P	81	E	82	D	83	F	84	A
Vk2		A		S		G		V		P		D		R		F		S		G		S		G		S		G		T		D		F		T		L		K		I		S		R		V		E		A		E		D		V		G
Vk3		A		T		G		V		P		A		R		F		S		G		S		G		S		G		T		D		F		T		L		T		I		S		S		L		E		P		E		D		F		A
Vk4		E		S		G		V		P		D		R		F		S		G		S		G		S		G		T		D		F		T		L		T		I		S		S		L		Q		A		E		D		V		A

framework 3		CDRIII																framework 4																																
Vk1	85	T	86	Y	87	Y	88	C	89	Q	90	Q	91	H	92	Y	93	T	94	T	95	P	96	P	97	T	98	F	99	G	100	Q	101	G	102	T	103	K	104	V	105	E	106	I	107	K	108	R	109	T
Vk2		V		Y		Y		C		Q		Q		H		Y		T		T		P		P		T		F		G		Q		G		T		K		V		E		I		K		R		T
Vk3		V		Y		Y		C		Q		Q		H		Y		T		T		P		P		T		F		G		Q		G		T		K		V		E		I		K		R		T
Vk4		V		Y		Y		C		Q		Q		H		Y		T		T		P		P		T		F		G		Q		G		T		K		V		E		I		K		R		T

[illegible][illegible]

[The following page contains musical notation.]

Figure 2B: VL lambda consensus sequences

framework 3	
	87
	86
	85
	84
	83
	82
	81
	80
	79
	78
	77
	76
	75
	74
	73
	72
	71
	70
	69
	68
	67
	66
	65
	64
	63
	62
	61
	60
	59
	58
VA1	V P D R F S G S K S G T S A S L A I T G L Q S E D E A D Y Y
VA2	V S N R F S G S K S G T A S L T I S G L Q A E D E A D Y Y
VA3	I P E R F S G S N S G N T A T L T I S G T Q A E D E A D Y Y

framework 4	
	107
	106
	105
	104
	103
	102
	101
	100
	99
	98
	97
	96
	95
	94
	93
	92
	91
	90
	89
	88
VA1	C Q Q H Y T T P P V F G G G T K L T V L G
VA2	C Q Q H Y T T P P V F G G G T K L T V L G
VA3	C Q Q H Y T T P P V F G G G T K L T V L G

Figure 2C: V heavy chain consensus sequences

framework 1		framework 2		CDR II	
VH1A	Q V Q L V Q S G A E V K K P G S S V K V S C K A S G G T F S	30	29	28	27
VH1B	Q V Q L V Q S G A E V K P G A S V K V S C K A S G Y T F T	26	25	24	23
VH2	Q V Q L K E S G P A L V K P T Q T L T L T C T F S G F S L S	22	21	20	19
VH3	E V Q L V E S G G L V Q P G S L R L S L T C T V S G G S I S	18	17	16	15
VH4	Q V Q L Q E S G P G L V K P S E T L S L T C T V S G G S I S	14	13	12	11
VH5	E V Q L V Q S G A E V K K P G E S L K I S C K G S G Y S F T	10	9	8	7
VH6	Q V Q L Q Q S G P G L V K P S Q T L S L T C A I S G D S V S	6	5	4	3
VH1A	S - - Y A I S W V R Q A P G Q G L E W M G G I I P - - I F G T A	57	56	55	54
VH1B	S - - Y Y M H W V R Q A P G Q G L E W M G W I N P - - N S G G T	53	52	51	50
VH2	T S G V G V G W I R Q P P G K A L E W L A L I D - - W D D D K	49	48	47	46
VH3	S - - Y A M S W V R Q A P G K G L E W V S A I S G - - S G G S T	45	44	43	42
VH4	S - - Y Y W S W I R Q P P G K G L E W I G Y I Y - - Y S G S T	41	40	39	38
VH5	S - - Y W I G W V R Q M P G K G L E W M G I I Y P - - G D S D T	37	36	35	34
VH6	S N S A A W N W I R Q S P G R G L E W L G R T Y Y R - S K W Y N	33	32	31	30

UNITED STATES OF AMERICA

THE UNIVERSITY OF CHICAGO

UNITED STATES OF AMERICA

Figure 3A: V kappa 1 (Vk1) gene sequence

```

.D I Q M T Q S P S S L S A S V G D
ECORV
~~~~~
BanII
~~~~~
GATATCCAGA TGACCCAGAG CCCGTCTAGC CTGAGCGCGA GCGTGGGTGA
CTATAGGTCT ACTGGGTCTC GGCAGATCG GACTCGCGCT CGCACCCACT

R V T I T C R A S Q G I S S Y L
PstI
~~~~~
TCGTGTGACC ATTACCTGCA GAGCGAGCCA GGGCATTAGC AGCTATCTGG
AGCACACTGG TAATGGACGT CTCGCTCGGT CCCGTAATCG TCGATAGACC

A W Y Q Q K P G K A P K L L I Y A
KpnI SexAI AseI
~~~~~
CGTGGTACCA GCAGAAACCA GGTAAAGCAC CGAAACTATT AATTATGCA
GCACCATGGT CGTCTTTGGT CCATTTCGTG GCTTTGATAA TTAAATACGT

A S S L Q S G V P S R F S G S
SandI BamHI
~~~~~
GCCAGCAGCT TGCAAAGCGG GGTCCCGTCC CGTTTAGCG GCTCTGGATC

```

CGGTCGTCGA ACGTTTCGCC CCAGGGCAGG GCAAATCGC CGAGACCTAG

G T D F T L T I S S L Q P E D F
Eco57I ~ ~ ~ ~ ~

BamHI

BbSI

~~~~~

CGGCACTGAT TTACCTGA CCATTAGCAG CCGCAACCT GAAGACTTTG  
GCCGTGACTA AAATGGGACT GGTAAATCGTC GGACGTTGGA CTTCTGAAAC

A T Y Y C Q Q H Y T T P P T F G Q  
MSCI

— 2  
2  
2  
2  
2

CGACCTATTA TTGCCAGCAG CATTATACCA CCCCGCCGAC CTTTGGCCAG  
GCTGGATAAT AACGGTCGTC GTAATATGGT GGGCGGCTG GAAACCGGTC

GTKEVET

BsiWI:

2  
2  
2  
2  
2

GGTACGAAG TTGAAATTAA ACGTACG  
CCATGCTTTC AACTTTAATT TGCATGC

**DECLASSIFICATION AUTHORITY**

[illegible]

Figure 3B: V kappa 2 [Nk2] gene sequence (continued)

|            |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
|------------|------------|------------|------------|------------|---|---|---|---|---|---|---|---|---|---|---|
| S          | G          | S          | G          | S          | G | T | D | F | T | L | K | I | S | R | V |
| BamHI      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| TAGCGGCTCT | GGATCCGGCA | CCGATTTTAC | CCTGAAAATT | AGCCGTGTGG |   |   |   |   |   |   |   |   |   |   |   |
| ATCGCCGAGA | CCTAGGCCGT | GGCTAAAATG | GGACTTTTAA | TCGGCACACC |   |   |   |   |   |   |   |   |   |   |   |
|            |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| E          | A          | E          | D          | V          | G | V | Y | Y | C | Q | Q | H | Y | T | P |
| Eco57I     |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| BbsI       |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| AAGCTGAAGA | CGTGGGCGTG | TATTATTGCC | AGCAGCATTA | TACCACCCCG |   |   |   |   |   |   |   |   |   |   |   |
| TTCGACTTCT | GCACCCGCAC | ATAATAACGG | TCGTCGTAAT | ATGGTGGGGC |   |   |   |   |   |   |   |   |   |   |   |
|            |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| P          | T          | F          | G          | Q          | G | T | K | V | E | I | K | R | T |   |   |
| MscI       |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| BsiWI      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| CCGACCTTTG | GCCAGGGTAC | GAAAGTTGAA | ATTAAACGTA | CG         |   |   |   |   |   |   |   |   |   |   |   |
| GGCTGGAAC  | CGGTCCCATG | CTTCAACT   | TAATTGCAT  | GC         |   |   |   |   |   |   |   |   |   |   |   |

[illegible]

GGCGCGAGCA GCCGTGCAAC TGGGTCCCG GCGGTTTA GCGCTCTGG



P E D  
Eco57I  
~~~~~

}
 }
 }
 }
 }

~~~~~  
ATCCGGCAG GATTTACCC TGACCATTAG CAGCCTGGAA CCTGAAGACT  
TAGGCCGTGC CTAAATGGG ACTGGTAATC GTCGGACCTT GGA CTCTGA

MSCI ~ ~ ~

TTGCGGTGTA TTATTGCCAG CAGCATTATA CCACCCCGCC GACCTTTGGC  
AACGCCACAT AATAACGGTC GTCGTAATAT GTGGGGCGG CTGGAACCCG

BsiWI ~~~~~

CAGGGTACGA AAGTTGAAA<sup>1</sup> TAAACGTACG  
GTCCCATGCT TTCAACTTTA ATTGCATGC

Figure 3D: V kappa 4 (Vx4) gene sequence

```

D I V M T Q S P D S L A V S L G E
EcoRV                               BanII
~~~~~
GATATCGTGA TGACCCAGAG CCCGATAGC CTGGCGGTGA GCCTGGGCGA
CTATAGCACT ACTGGGTCTC GGCCTATCG GACCGCCACT CGGACCCCGCT

R A T I N C R S S Q S V L Y S S
PstI
~~~~~
ACGTGCGACC ATTAAGTCA GAAGCAGCCA GAGCGTGCTG TATAGCAGCA
TGCACGCTGG TAATGACGT CTTCGTCGGT CTCGCACGAC ATATCGTCGT

N N K N Y L A W Y Q Q K P G Q P P
KpnI                               SexAI
~~~~~
ACAACAAAAA CTATCTGGCG TGGTACCAGC AGAAACCAGG TCAGCCGCCG
TGTTGTTTTT GATAGACCGC ACCATGGTCG TCCTTGGTCC AGTCGGCGGC

K L L I Y W A S T R E S G V P D R
AseI SmaI
~~~~~
AAACTATTAA TTTATTGGG ATCCACCCGT GAAAGCGGGG TCCCGGATCG
TTTGATAATT AAATAACCCG TAGGTGGGCA CTTTCGCCCC AGGGCCTAGC

```

Figure 3D: V kappa 4 (Vk4) gene sequence (continued)

|            |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
|------------|------------|------------|-------------|------------|---|---|---|---|---|---|---|---|---|---|---|
| F          | S          | G          | S           | G          | S | G | T | D | F | T | L | T | I | S | S |
| BamHI      |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| TTTTAGCGGC | TCTGGATCCG | GCACTGATT  | TACCCCTGACC | ATTTCGTCCC |   |   |   |   |   |   |   |   |   |   |   |
| AAAATCGCCG | AGACCTAGGC | CGTGACTAAA | ATGGGACTGG  | TAAAGCAGGG |   |   |   |   |   |   |   |   |   |   |   |
| L          | Q          | A          | E           | D          | V | A | V | Y | C | Q | Q | H | Y | T | T |
| Eco57I     |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| BbsI       |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| TGCAAGCTGA | AGACGTGGCG | GTGTATTATT | GCCAGCAGCA  | TTATACCACC |   |   |   |   |   |   |   |   |   |   |   |
| ACGTCGACT  | TCTGCACCGC | CACATAATAA | CGGTCGTCGT  | AATATGGTGG |   |   |   |   |   |   |   |   |   |   |   |
| P          | P          | T          | F           | G          | Q | G | T | K | V | E | I | K | R | T |   |
| MscI       |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| CCGCCGACCT | TTGGCCAGGG | TACGAAAGTT | GAAATTAAAC  | GTACG      |   |   |   |   |   |   |   |   |   |   |   |
| GGCGGCTGGA | AACCGGTCCC | ATGCTTTCAA | CTTTAATTG   | CATGC      |   |   |   |   |   |   |   |   |   |   |   |
| BsiWI      |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |             |            |   |   |   |   |   |   |   |   |   |   |   |

Figure 4A: V lambda 1 (Vλ1) gene sequence

Q S V L T Q P P S V S G A P G Q R  
SexAI

CAGAGCGTGC TGACCCAGCC GCCTTCAGTG AGTGGCGCAC CAGTCAGCG  
GTCTCGCAGC ACTGGGTCGG CGGAAGTCAC TCACCGCGTG GTCCAGTCGC  
Eco57I

V T I S C S G S S N I G S N Y  
BssSI

TGTGACCATC TCGTGTAGCG GCAGCAGCAG CAACATTGGC AGCAACTATG  
ACACTGGTAG AGCACATCGC CGTCGTCGTC GTTGTAACCG TCGTTGATAC

V S W Y Q Q L P G T A P K L L I Y  
KpnI XmaI BbeI

TGAGCTGGTA CCAGCAGTTG CCCGGGACGG CGCCGAAACT GCTGATTAT  
ACTCGACCAT GGTGTCACAC GGGCCCTGCC GCGGCTTTGA CGACTAAATA

D N N Q R P S G V P D R F S G S K  
Bsu36I BamHI

Figure 4A: V lambda 1 (Vλ1) gene sequence (continued)

GATAACAACC AGCGTCCCTC AGCGGTGCCG GATCGTTTAA GCGGATCCAA  
CTATTGTTGG TCGCAGGGAG TCCGCACGGC CTAGCAAAT CCGCTAGGTT

S G T S A S L A I T G L Q S E D  
BbsI

~~~~~  
AAGCGGCACC AGCGGAGCC TTGCGATTAC GGGCTGCAA AGCGAAGACG
TTCGCCGTGG TCGCGCTCGG AACGCTAATG CCCGGACGTT TCGCTTCTGC

E A D Y Y C Q Q H Y T T P P V F G
AAGCGGATTA TTATTGCCAG CAGCATTATA CCACCCCGCC TGTGTTTGGC
TTCGCCCTAAT AATAACGGTC GTCGTAATAT GGTGGGGCGG ACACAAACCG

G G T K L T V L G
HpaI MscI

~~~~~  
GGCGGCACGA AGTTAACCGT TCCTGGC  
CCGCCGTGCT TCAATTGGCA AGAACCG



Figure 4B: V lambda 2 (Vλ2) gene sequence (continued)

|                                  |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
|----------------------------------|---|---|---|---|---|---|---|---|---|-------|---|---|---|---|---|
| K                                | S | G | N | T | A | S | L | T | I | S     | G | L | Q | A | E |
| BamHI                            |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| ~~~~~                            |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| C                                | A | A | A | G | C | G | C | A | A | C     | C | G | C | G | A |
| GCTGACCAT TAGCGGCCTG CAAGCGGAAG  |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| G                                | T | T | T | C | G | C | C | G | T | T     | G | G | C | G | A |
| ATCGCCGGAC GTTCGCCCTC            |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| ~~~~~                            |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| D                                | E | A | D | Y | Y | C | Q | Q | H | Y     | T | T | P | P | V |
| F                                |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| BbsI                             |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| ~~~~~                            |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| A                                | C | G | A | A | G | C | G | G | A | T     | T | A | T | T | G |
| CAGCAGCATT ATACCACCCC GCCGTGTGTT |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| T                                | G | C | T | T | C | G | C | C | T | A     | A | T | A | A | C |
| GTCGTCGTAA TATGGTGGG CGGACACAAA  |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| ~~~~~                            |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| G                                | G | G | T | K | L | T | V | L | G |       |   |   |   |   |   |
| HpaI                             |   |   |   |   |   |   |   |   |   | MscI  |   |   |   |   |   |
| ~~~~~                            |   |   |   |   |   |   |   |   |   | ~~~~~ |   |   |   |   |   |
| G                                | G | G | C | G | G | C | A | A | G | T     | T | A | A | C | G |
| CGTCTTTGGC                       |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |
| C                                | C | G | C | C | G | C | C | G | T | T     | C | A | A | T | T |
| GCAAGAACCG                       |   |   |   |   |   |   |   |   |   |       |   |   |   |   |   |

[illegible]

# SexAI

Eco57I

**ISSI**

## Bbei





Figure 5A: V heavy chain 1A (VH1A) gene sequence

```

Q V Q L V Q S G A E V K K P G S S
MfeI
~~~~~
CAGGTGCAAT TGGTTCAGTC TGGCGCGGAA GTGAAAAAAC CGGGCAGCAG
GTCCACGTTA ACCAAGTCAG ACCGGCCCTT CACTTTTTTG GCCCGTCGTC

V K V S C K A S G G T F S S Y A
BspEI
~~~~~
CGTGAAAGTG AGCTGCAAAG CCTCCGGAGG CACTTTTAGC AGCTATGCCA
GCACTTTCAC TCGACGTTTC GGAGGCCTCC GTGAAAAATCG TCGATACGCT

I S W V R Q A P G Q G L E W M G G
BstXI XhoI
~~~~~
TTAGCTGGGT GCGCCAAGCC CCTGGGCAGG GTCTCGAGTG GATGGCGGC
AATCGACCCA CGCGGTTCCG GGACCCGTCC CAGAGCTCAC CTACCCGCCG

I I P I F G T A N Y A Q K F Q G R
ATTATCCGA TTTTGGCAC GCGAAGTAC GCGCAGAAGT TTCAGGGCCG
TAATAAGGCT AAAAACCCTG CCGCTTGATG CGCGTCTTCA AAGTCCCGGC

V T I T A D E S T S T A Y M E L
BstEII

```

SUBSTITUTE SHEET (RULE 26)

[illegible][illegible][illegible][illegible][illegible][illegible][illegible][illegible][illegible]

[illegible]

I N P N S G G T N Y A Q K F Q G R  
ATTAAACCGA ATAGCGGCGG CACGAAC TAC GCGAGAAGT TTCAGGGCCG  
TAATTGGGCT TATCGCCGCC GTGCTTGATG CGGCTCTCA AAGTCCCCGC

Figure 5B: V heavy chain 1B (VH1B) gene sequence (continued)

|            |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
|------------|------------|-------------|------------|------------|---|---|---|---|---|---|---|---|---|---|---|
| V          | T          | M           | T          | R          | D | T | S | I | S | T | A | Y | M | E | L |
| BstEII     |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| GGTGACCATG | ACCCGTGATA | CCAGCATTAG  | CACCGCGTAT | ATGGAAGTGA |   |   |   |   |   |   |   |   |   |   |   |
| CCACTGGTAC | TGGGCACTAT | GGTCGTAATC  | GTGGCGCATA | TACCTTGACT |   |   |   |   |   |   |   |   |   |   |   |
|            |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| S          | S          | L           | R          | S          | E | D | T | A | V | Y | Y | C | A | R | W |
| EagI       |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| GCAGCCTGCG | TAGCGAAGAT | ACGGCCGTGT  | ATTATTGCGC | GCGTTGGGGC |   |   |   |   |   |   |   |   |   |   |   |
| CGTCGGACGC | ATCGCTTCTA | TGCCGGCACCA | TAATAACGCG | CGCAACCCCG |   |   |   |   |   |   |   |   |   |   |   |
|            |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| G          | D          | G           | F          | Y          | A | M | D | Y | W | G | Q | G | T | L | V |
| StyI       |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| GGCGATGGCT | TTTATGCGAT | GGATTATTGG  | GGCCAAGGCA | CCCTGGTGAC |   |   |   |   |   |   |   |   |   |   |   |
| CCGCTACCGA | AAATACGCTA | CCTAATAACC  | CCGGTTCCGT | GGGACCACTG |   |   |   |   |   |   |   |   |   |   |   |
|            |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| V          | S          | S           |            |            |   |   |   |   |   |   |   |   |   |   |   |
| BlnI       |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| GGTAGCTCA  | G          |             |            |            |   |   |   |   |   |   |   |   |   |   |   |
| CCAATCGAGT | C          |             |            |            |   |   |   |   |   |   |   |   |   |   |   |

Figure 5C: V heavy chain 2 (VH2) gene sequence

```

Q V Q L K E S G P A L V K P T Q T
MfeI
~~~~~
CAGGTGCAAT TGAAGAAAG CGGCCCGGCC CTGGTGAAC CGACCCAAC
GTCCACGTTA ACTTCTTTC GCCGGGCCGG GACCACTTG GCTGGGTTTG

L T L T C T F S G F S L S T S G
BspEI
~~~~~
CCTGACCCTG ACCTGTACCT TTTCCGGATT TAGCCTGTCC ACGTCTGGCG
GGACTGGGAC TGGACATGGA AAAGGCCTAA ATCGGACAGG TGCAGACCGC

V G V G W I R Q P P G K A L E W L
BstXI XhoI
~~~~~
TTGGCGTGGG CTGGATTGCG CAGCCGCCCTG GAAAGCCCT CGAGTGGCTG
AACCGCACCC GACCTAAGCG GTCGGCGGAC CCTTCGGGA GCTCACCGAC

A L I D W D D D K Y Y S T S L K T
MluI
~~~~~
GCTCTGATTG ATTGGGATGA TGATAAGTAT TATAGCACCA GCCTGAAAC
CGAGACTAAC TAACCCTACT ACTATTCTATA ATATCGTGGT CGGACTTTTG

```

|                                                                                                                             |            |            |            |             |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
|-----------------------------------------------------------------------------------------------------------------------------|------------|------------|------------|-------------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|--|--|--|--|--|--|--|--|--|--|--|--|
| R                                                                                                                           | L          | T          | I          | S           | K | D | T | S | K | N | Q | V | V | L | T |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| <div style="display: flex; justify-content: space-between;"> <div> MluI<br/>~~~~~ </div> <div> NspV<br/>~~~~~ </div> </div> |            |            |            |             |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| GCGTCTGACC                                                                                                                  | ATTAGCAAAG | ATACTTCGAA | AAATCAGGTG | GTGCTGACTA  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| CGCAGACTGG                                                                                                                  | TAATCGTTTC | TATGAAGCTT | TTTAGTCCAC | CACGACTGAT  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| M                                                                                                                           | T          | N          | M          | D           | P | V | D | T | A | T | Y | Y | C | A | R | W |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| <div style="display: flex; justify-content: space-between;"> <div>BssHII</div> <div>~~~~~</div> </div>                      |            |            |            |             |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| TGACCAACAT                                                                                                                  | GGACCCGGTG | GATACGGCCA | CCTATTATTG | CGCGCGTTGG  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| ACTGTTGTA                                                                                                                   | CCTGGGCCAC | CTATGCCGGT | GGATAATAAC | CGCGCAACC   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| G                                                                                                                           | G          | D          | G          | F           | Y | A | M | D | Y | W | G | Q | G | T | L | V |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| <div style="display: flex; justify-content: space-between;"> <div>StyI</div> <div>~~~~~</div> </div>                        |            |            |            |             |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| GGCGGCGATG                                                                                                                  | GCTTTATGC  | GATGGATTAT | TGGGGCCAAG | GCACCCCTGGT |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| CCGCCGCTAC                                                                                                                  | CGAAATACG  | CTACCTAATA | ACCCCGGTC  | CGTGGGACCA  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| <div style="display: flex; justify-content: space-between;"> <div>T</div></div>                                             |            |            |            |             |   |   |   |   |   |   |   |   |   |   |   | V | S | S |  |  |  |  |  |  |  |  |  |  |  |  |
| <div style="display: flex; justify-content: space-between;"> <div>BlpI</div> <div>~~~~~</div> </div>                        |            |            |            |             |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| GACGGTTAGC                                                                                                                  | TCAG       |            |            |             |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |
| CTGCCAATCG                                                                                                                  | AGTC       |            |            |             |   |   |   |   |   |   |   |   |   |   |   |   |   |   |  |  |  |  |  |  |  |  |  |  |  |  |

Figure 5D: V heavy chain 3 (VH3) gene sequence

```

E V Q L V E S G G G L V Q P G G S
MfeI
~~~~~
GAAGTGCAAT TGGTGGAAG CGGCGGCGGC CTGGTGCAAC CGGCGGCGCAG
CTTCACGTTA ACCACCTTC GCCGCCGCCG GACCACGTTG GCCCGCCGTC

L R L S C A A S G F T F S S Y A
BspEI
~~~~~
CCTGCGTCTG AGCTGCGCGG CCTCCGGATT TACCTTTAGC AGCTATGCGA
GGACGCAGAC TCGACGCGCC GGAGGCCTAA ATGGAAATCG TCGATACGCT

M S W V R Q A P G K G L E W V S A
BstXI
~~~~~
XhoI
~~~~~
TGAGCTGGGT GCGCCAAGCC CCTGGGAAGG GTCTCGAGTG GGTGAGCGCG
ACTCGACCCA CGCGGTTCGG GGACCCCTCC CAGAGCTCAC CCACTCGCGC

I S G S G G S T Y Y A D S V K G R
ATTAGCGGTA GCGCGGCGCAG CACCTATTAT GCGGATAGCG TGAAGGCCCG
TAATCGCCAT CGCCGCCGTC GTGGATAATA CGCCTATCGC ACTTCCCGGC

```



2010-01-01

[illegible]

Figure 5E: V heavy chain 4 (VH4) gene sequence

```

Q V Q L Q E S G P G L V K P S E T
MfeI
~~~~~
CAGGTGCAAT TGCAAGAAAG TGGTCCGGGC CTGGTGAAC CGAGCGAAAC
GTCCACGTTA ACGTTCCTTC ACCAGGCCCG GACCACTTG GCTCGCTTG

L S L T C T V S G G S I S S Y Y
BspEI
~~~~~
CCTGAGCCTG ACCTGCACCG TTTCCGGAGG CAGCATTAGC AGCTATTATT
GGA CTGGAC TGGACGTGGC AAAGCCTCC GTCGTAATCG TCGATAATAA

W S W I R Q P P G K G L E W I G Y
BstXI
~~~~~
XhoI
~~~~~
GGAGCTGGAT TCGCCAGCCG CCTGGGAAGG GTCTCGAGTG GATTGGCTAT
CCTCGACCCTA AGCGGTCGGC GGACCCTTCC CAGAGCTCAC CTAACCGATA

I Y Y S G S T N Y N P S L K S R V
BstEII
~~~~~
ATTATTATA GCGGCAGCAC CAACTATAAT CCGAGCCTGA AAAGCCGGGT
TAAATAATAT CGCCGTCGTG GTTGATATTA GGCTCGGACT TTTCGGCCCA

```

Figure 5E: V heavy chain 4 (VH4) gene sequence (continued)

|        |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
|--------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| T      | I | S | V | D | T | S | K | N | Q | F | S | L | K | L | S |   |   |   |
| BstEII |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| NspV   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| G      | A | C | C | A | T | T | A | C | T | T | G | A | A | A | A | C | C | A |
| C      | T | G | G | T | A | A | T | A | T | G | A | C | G | A | C | A | A | A |
| ~~~~~  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| S      | V | T | A | A | D | T | A | V | Y | Y | C | A | R | W | G | G |   |   |
| EagI   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| G      | C | G | T | G | A | C | G | G | A | T | A | C | G | G | T | A | T |   |
| C      | G | C | A | T | G | C | C | G | T | A | T | G | C | G | C | G | C |   |
| ~~~~~  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| D      | G | F | Y | A | M | D | Y | W | G | Q | G | T | L | V | T | V |   |   |
| StyI   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| G      | A | T | G | G | C | T | T | T | A | T | G | G | A | T | G | G | C |   |
| C      | T | A | C | C | G | A | A | A | A | A | A | A | A | A | A | A | A |   |
| ~~~~~  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| S      | S |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| BlnI   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |
| T      | A | G | C | T | C | A | G |   |   |   |   |   |   |   |   |   |   |   |
| A      | T | C | G | A | G | T | C |   |   |   |   |   |   |   |   |   |   |   |

Figure 5F: V heavy chain 5 (VH5) gene sequence

```

E V Q L V Q S G A E V K K P G E S
MfeI
~~~~~
GAAGTGCAAT TGGTTCAGAG CGGCGCGGAA GTGAAAAAAC CGGGCGAAAG
CTTCACGTTA ACCAAGTCTC GCCGGCCTT CACTTTTTCG GCCCGCTTTC

L K I S C K G S G Y S F T S Y W
BspEI
~~~~~
CCTGAAAATT AGCTGCAAAG GTTCCGGGATA TTCCTTTACG AGCTATTGGA
GGACTTTTAA TCGACGTTTC CAAGGCCTAT AAGGAAATGC TCGATAACCT

I G W V R Q M P G K G L E W M G I
BstXI
~~~~~
XhoI
~~~~~
TTGGCTGGGT GCGCCAGATG CCTGGGAAGG GTCTCGAGTG GATGGGCATT
AACCGACCCA CGCGGTCTAC GGACCCCTTC CAGAGCTCAC CTACCCGTAA

I Y P G D S D T R Y S P S F Q G Q
ATTATCCGG GCGATAGCGA TACCCGTTAT TCTCCGAGCT TTCAGGGCCA
TAAATAGGCC CGCTATCGCT ATGGGCAATA AGAGGCTCGA AAGTCCCCGGT

```

Figure 5F: V heavy chain 5 (VH5) gene sequence (continued)

|            |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
|------------|------------|------------|------------|------------|---|---|---|---|---|---|---|---|---|---|---|
| V          | T          | I          | S          | A          | D | K | S | I | S | T | A | Y | L | Q | W |
| BstEII     |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| GGTGACCAT  | AGCGCGGATA | AAAGCATTAG | CACCGCGTAT | CTTCAATGGA |   |   |   |   |   |   |   |   |   |   |   |
| CCACTGGTAA | TCGCGCCTAT | TTTCGTAATC | GTGGCGCATA | GAAGTTACCT |   |   |   |   |   |   |   |   |   |   |   |
| S          | S          | L          | K          | A          | S | D | T | A | M | Y | Y | C | A | R | W |
| G          |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| BssHII     |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| GCAGCCTGAA | AGCGAGCGAT | ACGGCCATGT | ATTATTGCGC | CGTTGGGGC  |   |   |   |   |   |   |   |   |   |   |   |
| CGTCGGACTT | TCGCTCGCTA | TGCCGGTACA | TAATAACGCG | CGCAACCCCG |   |   |   |   |   |   |   |   |   |   |   |
| G          | D          | G          | F          | Y          | A | M | D | Y | W | G | Q | G | T | L | V |
| T          |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| StyI       |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| GGCGATGGCT | TTTATGCGAT | GGATTATTGG | GGCCAAGGCA | CCCTGGTGAC |   |   |   |   |   |   |   |   |   |   |   |
| CCGCTACCGA | AAATACGCTA | CCTAATAACC | CCGTTCCGT  | GGGACCACTG |   |   |   |   |   |   |   |   |   |   |   |
| V          | S          | S          |            |            |   |   |   |   |   |   |   |   |   |   |   |
| BlpI       |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| ~~~~~      |            |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| GGTAGCTCA  | G          |            |            |            |   |   |   |   |   |   |   |   |   |   |   |
| CCAATCGAGT | C          |            |            |            |   |   |   |   |   |   |   |   |   |   |   |

Figure 5G: V heavy chain 6 (VH6) gene sequence

```

Q V Q L Q Q Q S G P G L V K P S Q T
MfeI
~~~~~
CAGGTGCAAT TGCAACAGTC TGGTCCGGGC CTGGTGAAC CGAGCCAAAC
GTCCACGTTA ACGTTGTCAG ACCAGGCCCG GACCACTTG GTCGGTTTG

L S L T C A I S G D S V S S N S
BspEI
~~~~~
CCTGAGCCTG ACCTGTGCGA TTTCCGGAGA TAGCGTGAGC AGCAACAGCG
GGA CTGGAC TGGACACGCT AAAGGCCTCT ATCGCACTCG TCGTTGTGCG

A A W N W I R Q S P G R G L E W L
BstXI XhoI
~~~~~
CGGCGTGGAA CTGGATTGCG CAGTCTCCTG GCGTGCCCT CGAGTGGCTG
GCCGCACCTT GACCTAAGCG GTCAGAGGAC CCGCACCGGA GTCACCCGAC

G R T Y Y R S K W Y N D Y A V S V
GGCCGTACCT ATTATCGTAG CAAATGGTAT AACGATTATG CCGTGAGCGT
CCGGCATGGA TAATAGCATC GTTTACCATA TTGCTAATAC GCCACTCGCA

```

Figure 5G: V heavy chain 6 (VH6) gene sequence (continued)

|   |   |   |   |   |   |   |   |   |   |        |   |   |   |   |   |
|---|---|---|---|---|---|---|---|---|---|--------|---|---|---|---|---|
| K | S | R | I | T | I | N | P | D | T | S      | K | N | Q | F | S |
|   |   |   |   |   |   |   |   |   |   | NspV   |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | ~~~~~  |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | BsaBI  |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | ~~~~~  |   |   |   |   |   |
| G | A | A | A | G | C | C | G | G | A | T      | T | A | C | C | A |
| T | C | A | A | A | A | A | A | A | A | C      | C | C | C | G | G |
| C | T | T | T | C | G | G | C | C | T | A      | T | T | T | T | G |
| T | G | G | C | C | C | T | A | T | G | A      | A | G | C | T | T |
| T | T | T | T | T | T | T | T | T | T | T      | T | T | T | T | T |
| L | Q | L | N | S | V | T | P | E | D | T      | A | V | Y | C | A |
|   |   |   |   |   |   |   |   |   |   | EagI   |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | ~~~~~  |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | BssHII |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | ~~~~~  |   |   |   |   |   |
| T | G | C | A | A | C | T | G | A | A | G      | A | T | A | C | C |
| C | C | G | G | A | A | G | A | T | A | C      | C | G | T | G | G |
| A | C | G | T | T | G | A | C | T | T | T      | A | T | T | T | G |
| G | C | C | G | C | C | T | T | C | T | A      | T | A | T | A | C |
| G | C | C | G | C | C | T | A | T | A | C      | C | A | T | A | C |
| R | W | G | G | D | G | F | Y | A | M | D      | Y | W | G | Q | T |
|   |   |   |   |   |   |   |   |   |   | StyI   |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | ~~~~~  |   |   |   |   |   |
| C | G | T | T | G | G | G | C | G | T | T      | A | T | T | G | G |
| G | C | A | A | C | C | C | G | C | T | A      | T | A | T | T | G |
| G | C | A | A | C | C | C | G | C | T | A      | T | A | T | T | G |
| L | V | T | V | S | S |   |   |   |   |        |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | BlnI   |   |   |   |   |   |
|   |   |   |   |   |   |   |   |   |   | ~~~~~  |   |   |   |   |   |
| C | C | T | G | T | G | A | C | G | T | A      | G | C | T | C | A |
| G | G | A | C | C | A | C | T | G | C | A      | T | C | G | A | G |

Figure 6: oligonucleotides for gene synthesis

O1K1 5' - GAATGCATACGCTGATATCCAGATGACCCAGAG-  
CCCGTCTAGCCTGAGC -3'

O1K2 5' - CGCTCTGCAGGTAATGGTCACACGATCACCCAC-  
GCTCGCGCTCAGGCTAGACGGGC -3'

O1K3 5' - GACCATTACCTGCAGAGCGAGCCAGGGCATTAG-  
CAGCTATCTGGCGTGGTACCAGCAG -3'

O1K4 5' - CTTTGCAAGCTGCTGGCTGCATAAATTAATAGT-  
TTCGGTGCTTTACCTGGTTTCTGCTGGTACCACGCCAG -3'

O1K5 5' - CAGCCAGCAGCTTGCAAAGCGGGGTCCCGTCCC-  
GTTTTAGCGGCTCTGGATCCGGCACTGATTTTAC -3'

O1K6 5' - GATAATAGGTCGCAAAGTCTTCAGGTTGCAGGC-  
TGCTAATGGTCAGGGTAAAATCAGTGCCGGATCC -3'

O2K1 5' - CGATATCGTGATGACCCAGAGCCCACTGAGCCT-  
GCCAGTGACTCCGGGGCGAGCC -3'

O2K2 5' - GCCGTTGCTATGCAGCAGGCTTTGGCTGCTTCT-  
GCAGCTAATGCTCGCAGGCTCGCCCCGGAGTCAC -3'

O2K3 5' - CTGCTGCATAGCAACGGCTATAACTATCTGGAT-  
TGGTACCTTCAAAAACCAGGTCAAAGCCC -3'

O2K4 5' - CGATCCGGGACCCCACTGGCACGGTTGCTGCCC-  
AGATAAATTAATAGCTGCGGGCTTTGACCTGGTTTTTG -3'

O2K5 5' - AGTGGGGTCCCGGATCGTTTTAGCGGCTCTGGA-  
TCCGGCACCGATTTTACCCTGAAAATTAGCCGTGTG -3'

O2K6 5' - CCATGCAATAATACACGCCCACGTCTTCAGCTT-  
CCACACGGCTAATTTTCAGGG -3'

O3K1 5' - GAATGCATACGCTGATATCGTGCTGACCCAGAG-  
CCCGG -3'

O3K2 5' - CGCTCTGCAGCTCAGGGTCGCACGTTGCCCCGG-  
AGACAGGCTCAGGGTCGCCGGGCTCTGGGTCAGC -3'

O3K3 5' - CCCTGAGCTGCAGAGCGAGCCAGAGCGTGAGCA-  
GCAGCTATCTGGCGTGGTACCAG -3'

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Figure 6: (continued)

**O3K4** 5' - GCACGGCTGCTCGCGCCATAAATTAATAGACGC-  
GGTGCTTGACCTGGTTTCTGCTGGTACCACGCCAGATAG -3'

**O3K5** 5' - GCGCGAGCAGCCGTGCAACTGGGGTCCCGGCGC-  
GTTTTAGCGGCTCTGGATCCGGCACGGATTTTAC -3'

**O3K6** 5' - GATAATACACCGCAAAGTCTTCAGGTTCCAGGC-  
TGCTAATGGTCAGGGTAAAATCCGTGCCGGATC -3'

**O4K1** 5' - GAATGCATACGCTGATATCGTGATGACCCAGAG-  
CCCGGATAGCCTGGCG -3'

**O4K2** 5' - GCTTCTGCAGTTAATGGTCGCACGTTTCGCCCAG-  
GCTCACCGCCAGGCTATCCGGGC -3'

**O4K3** 5' - CGACCATTAAGTGCAGAAGCAGCCAGAGCGTGC-  
TGTATAGCAGCAACAACAAAACTATCTGGCGTGGTACCAG -  
3'

**O4K4** 5' - GATGCCCAATAAATTAATAGTTTCGGCGGCTGA-  
CCTGGTTTCTGCTGGTACCACGCCAGATAG -3'

**O4K5** 5' - AAATATTAATTTATTGGGCATCCACCCGTGAA-  
AGCGGGGTCCCGGATCGTTTTAGCGGCTCTGGATCCGGCAC-  
3'

**O4K6** 5' - GATAATACACCGCCACGTCTTCAGCTTGCAGGG-  
ACGAAATGGTCAGGGTAAAATCAGTGCCGGATCCAGAGCC -  
3'

**O1L1** 5' - GAATGCATACGCTCAGAGCGTGCTGACCCAGCC-  
GCCTTCAGTGAGTGG -3'

**O1L2** 5' - CAATGTTGCTGCTGCTGCCGCTACACGAGATGG-  
TCACACGCTGACCTGGTGCGCCACTCACTGAAGGCGGC -3'

**O1L3** 5' - GGCAGCAGCAGCAACATTGGCAGCAACTATGTG-  
AGCTGGTACCAGCAGTTGCCCCGGGAC -3'

**O1L4** 5' - CCGGCACGCCTGAGGGACGCTGGTTGTTATCAT-  
AAATCAGCAGTTTCGGCGCCGTCCCGGGCAACTGC -3'

**O1L5** 5' - CCCTCAGGCGTGCCGGATCGTTTTAGCGGATCC-  
AAAAGCGGCACCAGCGCGAGCCTTGCG -3'

Figure 6: (continued)

O1L6 5' - CCGCTTCGTCTTCGCTTTGCAGGCCCCGTAATCG-  
CAAGGCTCGCGCTGG -3'

O2L1 5' - GAATGCATACGCTCAGAGCGCACTGACCCAGCC-  
AGCTTCAGTGAGCGGC -3'

O2L2 5' - CGCTGCTAGTACCCGTACACGAGATGGTAATGC-  
TCTGACCTGGTGAGCCGCTCACTGAAGCTGG -3'

O2L3 5' - GTACGGGTACTAGCAGCGATGTGGGCGGCTATA-  
ACTATGTGAGCTGGTACCAGCAGCATCCCCG -3'

O2L4 5' - CGCCTGAGGGACGGTTGCTCACATCATAAATCA-  
TCAGTTTCGGCGCCTTCCCCGGGATGCTGCTGGTAC -3'

O2L5 5' - CAACCGTCCCTCAGGCGTGAGCAACCGTTTTAG-  
CGGATCCAAAAGCGGCAACACCGCGAGCC -3'

O2L6 5' - CCGCTTCGTCTTCCGCTTGCAGGCCGCTAATGG-  
TCAGGCTCGCGGTGTTGCCG -3'

O3L1 5' - GAATGCATACGCTAGCTATGAACTGACCCAGCC-  
GCCTTCAGTGAGCG -3'

O3L2 5' - CGCCCAGCGCATCGCCGCTACACGAGATACGCG-  
CGGTCTGACCTGGTGCAACGCTCACTGAAGGCGGC -3'

O3L3 5' - GGCGATGCGCTGGGCGATAAATACGCGAGCTGG-  
TACCAGCAGAAACCCGGGCAGGCGC -3'

O3L4 5' - GCGTTCCGGGATGCCTGAGGGACGGTCAGAATC-  
ATCATAAATCACCAGAACTGGCGCCTGCCCCGGGTTTC -3'

O3L5 5' - CAGGCATCCCGGAACGCTTTAGCGGATCCAACA-  
GCGGCAACACCGCGACCCTGACCATTAGCGG -3'

O3L6 5' - CCGCTTCGTCTTCCGCCTGAGTGCCGCTAATGG-  
TCAGGGTC -3'

O1246H1 5' - GCTCTTCACCCCTGTTACCAAAGCCCAG-  
GTGCAATTG -3'

O1AH2 5' - GGCTTTGCAGCTCACTTTCACGCTGCTGCCCCGG-  
TTTTTTCACTTCCGCGCCAGACTGAACCAATTGCACCTGGGC-  
TTTG -3'

Figure 6: (continued)

**O1AH3** 5' - GAAAGTGAGCTGCAAAGCCTCCGGAGGGCACTTT-  
TAGCAGCTATGCGATTAGCTGGGTGCGCCAAGCCCCTGGGCAG  
GGTC -3'

**O1AH4** 5' - GCCCTGAAACTTCTGCGCGTAGTTCGCCGTGCC-  
AAAAATCGGAATAATGCCGCCCATCCACTCGAGACCCTGCCC-  
AGGGGC -3'

**O1AH5** 5' - GCGCAGAAGTTTCAGGGCCGGGTGACCATTACC-  
GCGGATGAAAGCACCAGCACC GCGTATATGGAAGT GAGCAGCC  
TGCG -3'

**O1ABH6** 5' - GCGCGCAATAATACACGGCCGTATCTTCGCT-  
ACGCAGGCTGCTCAGTTCC -3'

**O1BH2** 5' - GGCTTTGCAGCTCACTTTCACGCTCGCGCCCGG-  
TTTTTTCCTTCCGCGCCGCTCTGAACCAATTGCACCTGGGC-  
TTTG -3'

**O1BH3** 5' - GAAAGTGAGCTGCAAAGCCTCCGGATATACCTT-  
TACCAGCTATTATATGCACTGGGTCCGCCAAGCCCCTGGGCAG  
GGTC -3'

**O1BH4** 5' - GCCCTGAAACTTCTGCGCGTAGTTCGTGCCGCC-  
GCTATTCGGGTTAATCCAGCCCATCCACTCGAGACCCTGCCA  
GGGGC -3'

**O1BH5** 5' - GCGCAGAAGTTTCAGGGCCGGGTGACCATGACC-  
CGTGATACCAGCATTAGCACCGCGTATATGGAAGT GAGCAGCC  
TGCG -3'

**O2H2** 5' - GGTACAGGTCAGGGTCAGGGTTTGGGTCGGTTT-  
CACCAGGGCCGGGCGCTTTCTTTCAATTGCACCTGGGCTTTG  
-3'

**O2H3** 5' - CTGACCCTGACCTGTACCTTTTCCGGATTTAGC-  
CTGTCCACGTCTGGCGTTGGCGTGGGCTGGATTGCGCCAGCCGC  
CTGGGAAAG -3'

**O2H4** 5' - GCGTTTTTCAGGCTGGTGCTATAATACTTATCAT-  
CATCCCAATCAATCAGAGCCAGCCACTCGAGGGCTTTCCCAGG  
CGGCTGG -3'

Figure 6: (continued)

**O2H5** 5' - GCACCAGCCTGAAAACGCGTCTGACCATTAGCA-  
AAGATACTTCGAAAAATCAGGTGGTGCTGACTATGACCAACAT  
GG -3'

**O2H6** 5' - GCGCGCAATAATAGGTGGCCGTATCCACCGGGT-  
CCATGTTGGTCATAGTCAGC -3'

**O3H1** 5' - CGAAGTGCAATTGGTGGAAAGCGGCGGCGGCCT-  
GGTGCAACCGGGCGGCAG -3'

**O3H2** 5' - CATAGCTGCTAAAGGTAAATCCGGAGGCCGCGC-  
AGCTCAGACGCAGGCTGCCGCCCGGTTGCAC -3'

**O3H3** 5' - GATTTACCTTTAGCAGCTATGCGATGAGCTGGG-  
TGCGCCAAGCCCCCTGGGAAGGGTCTCGAGTGGGTGAG -3'

**O3H4** 5' - GGCCTTTCACGCTATCCGCATAATAGGTGCTGC-  
CGCCGCTACCGCTAATCGCGCTCACCCACTCGAGACCC -3'

**O3H5** 5' - CGGATAGCGTGAAAGGCCGTTTTACCATTTTAC-  
GTGATAATTCGAAAACACCCTGTATCTGCAAATGAACAG-3'

**O3H6** 5' - CACGCGCGCAATAATACACGGCCGTATCTTCCG-  
CACGCAGGCTGTTTCATTTGCAGATACAGG -3'

**O4H2** 5' - GGTCAGGCTCAGGGTTTCGCTCCTTTCACCAG-  
GCCCCGACCCTTTCTTGCAATTGCACCTGGGCTTTG -3'

**O4H3** 5' - GAAACCCTGAGCCTGACCTGCACCGTTTCCGGA-  
GGCAGCATTAGCAGCTATTATTGGAGCTGGATTCGCCAGCCGC  
-3'

**O4H4** 5' - GATTATAGTTGGTGCTGCCGCTATAATAAATAT-  
AGCCAATCCACTCGAGACCCTTCCCAGGCGGCTGGCGAATCCA  
G -3'

**O4H5** 5' - CGGCAGCACCAACTATAATCCGAGCCTGAAAAG-  
CCGGGTGACCATTAGCGTTGATACTTCGAAAACAGTTTAGC  
CTG -3'

**O4H6** 5' - GCGCGCAATAATACACGGCCGTATCCGCCGCCG-  
TCACGCTGCTCAGTTTCAGGCTAAACTGGTTTTTTCG -3'

Figure 6: (continued)

**O5H1** 5' - GCTCTTCACCCCTGTTACCAAAGCCGAAGTGCA-  
ATTG -3'

**O5H2** 5' - CCTTTGCAGCTAATTTTCAGGCTTTCGCCCCGGT-  
TTTTTCACTTCCGCGCCGCTCTGAACCAATTGCACTTCGGCTT  
TGG -3'

**O5H3** 5' - CCTGAAAATTAGCTGCAAAGGTTCCGGATATTC-  
CTTTACGAGCTATTGGATTGGCTGGGTGCGCCAGATGCCTGG  
-3'

**O5H4** 5' - CGGAGAATAACGGGTATCGCTATCGCCCCGGATA-  
AATAATGCCCATCCACTCGAGACCCTTCCCAGGCATCTGGCGC  
AC -3'

**O5H5** 5' - CGATACCCGTTATTCTCCGAGCTTTCAGGGCCA-  
GGTGACCATTAGCGCGGATAAAAGCATTAGCACCGCGTATCTT  
C -3'

**O5H6** 5' - GCGCGCAATAATACATGGCCGTATCGCTCGCTT-  
TCAGGCTGCTCCATTGAAGATACGCGGTGCTAATG -3'

**O6H2** 5' - GAAATCGCACAGGTCAGGCTCAGGGTTTGGCTC-  
GGTTTCACCAGGCCCGGACCAGACTGTGCAATTGCACCTGG-  
GCTTTG -3'

**O6H3** 5' - GCCTGACCTGTGCGATTTCCGGAGATAGCGTGA-  
GCAGCAACAGCGCGGCGTGGAAGTGGATTGCGCCAGTCTCCTGG  
GCG -3'

**O6H4** 5' - CACCGCATAATCGTTATAACCATTTGCTACGATA-  
ATAGGTACGGCCCAGCCACTCGAGGCCACGCCCAGGAGACTG-  
GCG -3'

**O6H5** 5' - GGTATAACGATTATGCGGTGAGCGTGAAAAGCC-  
GGATTACCATCAACCCGGATACTTCGAAAAACCAGTTTAGCCT  
GC -3'

**O6H6** 5' - GCGCGCAATAATACACGGCCGTATCTTCCGGGG-  
TCACGCTGTTTCAGTTGCAGGCTAAACTGGTTTTTC -3'

**OCLK1** 5' - GGCTGAAGACGTGGGCGTGTATTATTGCCAGCA-  
GCATTATACCACCCCGGACCTTTGGCCAGGGTAC -3'

Figure 6: (continued)

OCLK2 5' - GCGGAAAAATAAACACGCTCGGAGCAGCCACCG-  
TACGTTTAATTTCAACTTTCGTACCCTGGCCAAAGGTC -3'  
OCLK3 5' - GAGCGTGTTTATTTTCCGCCGAGCGATGAACA-  
ACTGAAAAGCGGCACGGCGAGCGTGCGTGTGCCTGCTG -3'  
OCLK4 5' - CAGCGCGTTGTCTACTTTCCACTGAACTTTCGC-  
TTCACGCGGATAAAAGTTGTTTCAGCAGGCACACCACGC -3'  
OCLK5 5' - GAAAGTAGACAACGCGCTGCAAAGCGGCAACAG-  
CCAGGAAAGCGTGACCGAACAGGATAGCAAAGATAG -3'  
OCLK6 5' - GTTTTTCATAATCCGCTTTGCTCAGGGTCAGGG-  
TGCTGCTCAGAGAATAGGTGCTATCTTTGCTATCCTGTTCG -  
3'  
OCLK7 5' - GCAAAGCGGATTATGAAAAACATAAAGTGTATG-  
CGTGCGAAGTGACCCATCAAGGTCTGAGCAGCCCGGTG -3'  
OCLK8 5' - GGCATGCTTATCAGGCCTCGCCACGATTAAAAG-  
ATTTAGTCACCGGGCTGCTCAGAC -3'  
OCH1 5' - GCGGTCTAGAGGCCAAGGCACCCTGGTGACGGT-  
TAGCTCAGCGTCGAC -3'  
OCH2 5' - GTGCTTTTGTGCTCGGAGCCAGCGGAAACACG-  
CTTGACCTTTGGTTCGACGCTGAGCTAACC -3'  
OCH3 5' - CTCCGAGCAGCAAAAGCACCAGCGGCGGCACGG-  
CTGCCCTGGGCTGCCTGGTTAAAGATTATTTC -3'  
OCH4 5' - CTGGTCAGCGCCCCGCTGTTCCAGCTCACGGTG-  
ACTGGTTCCGGGAAATAATCTTTAACCAGGCA -3'  
OCH5 5' - AGCGGGGCGCTGACCAGCGGCGTGCATACCTTT-  
CCGGCGGTGCTGCAAAGCAGCGGCCTG -3'  
OCH6 5' - GTGCCTAAGCTGCTGCTCGGCACGGTCACAACG-  
CTGCTCAGGCTATACAGGCCGCTGCTTTGCAG -3'  
OCH7 5' - GAGCAGCAGCTTAGGCACTCAGACCTATATTG-  
CAACGTGAACCATAAACCAGCAACACC -3'  
OCH8 5' - GCGCGAATTCGCTTTTTCGGTTCCACTTTTTTAT-  
CCTTTTGGTGTTGCTCGGTTTATGG -3'

V V A A A P S V F I F P P S D E Q Q

BSiWI

CGTACGGTGG CTGCTCCGAG CGTGTTATT TTTCGCCCA GCGATGAACA  
GCATGCCACC GACGAGGCTC GCACAAATAA AAAGGCGGCT CGCTACTTGT

|            |             |             |            |             |   |   |   |   |   |   |   |   |   |
|------------|-------------|-------------|------------|-------------|---|---|---|---|---|---|---|---|---|
| L          | K           | S           | G          | T           | A | S | V | C | L | L | N | F | Y |
| ACTGAAAGC  | GGCACGGCGA  | GCCTGGTGTG  | CCTGCTGAAC | AACTTTATC   |   |   |   |   |   |   |   |   |   |
| TGACTTTTCG | CCGTGCCCGCT | CGCACCCACAC | GGACGACTTG | TTGAAAAATAG |   |   |   |   |   |   |   |   |   |

P R E A K V Q W K V D N A L Q S G  
CGCGTGAAGC GAAAGTTCAG TGGAAAGTAG ACAACGCGCT GCAAAGCGGC  
GCGCACTTCG CTTCAAGTC ACCTTTCATC TGTTGCGCGA CGTTTCGCCG

N S Q E S V T E Q D S K D S T Y S  
AACAGCCAGG AAGCGTGAC CGAACAGGAT AGCAAGATA GCACCTATTC  
TTGTCGGTCC TTTCGCAC TGCTTGCTTA TCGTTTCTAT CGTGGATAAG

L S S T L T L S K A D Y E K H K  
TCTGAGCAGC ACCCTGACCC TGAGCAAGC GGATTATGAA AAACATAAAG  
AGACTCGTCG TGGGACTGGG ACTCGTTTCG CCTAATACTT TTTGTATTTT

Figure 7A: sequence of the synthetic Cx gene segment (continued)

V Y A C E V T H Q G L S S P V T K  
 TGTATGCGTG CGAAGTGACC CATCAAGGTC TGAGCAGCCC GGTGACTAAA  
 ACATACGCAC GCTTCACTGG GTAGTTCCAG ACTCGTCGGG CCACTGATTT

S F N R G E A \*

StuI SphI

~~~~~  
 TCTTTTAAATC GTGGCGAGGC CTGATAAGCA TGC  
 AGAAAATTAG CACCGCTCCG GACTATTCGT ACG



**WILLIAMSON**

[illegible]

[illegible]

|           |            |         |       |       |       |   |         |
|-----------|------------|---------|-------|-------|-------|---|---------|
| E         | P          | K       | S     | E     | F     | * | HindIII |
| AACCGAAAG | CGAATCTGA  | TAAGCTT | ~~~~~ | ECORI | ~~~~~ |   |         |
| TTGGCTTTC | GCTTAAGACT | ATTCGAA |       |       |       |   |         |

0011270-13006160

Figure 7C: functional map and sequence of module 24 comprising the synthetic Cλ gene segment (huCL lambda)

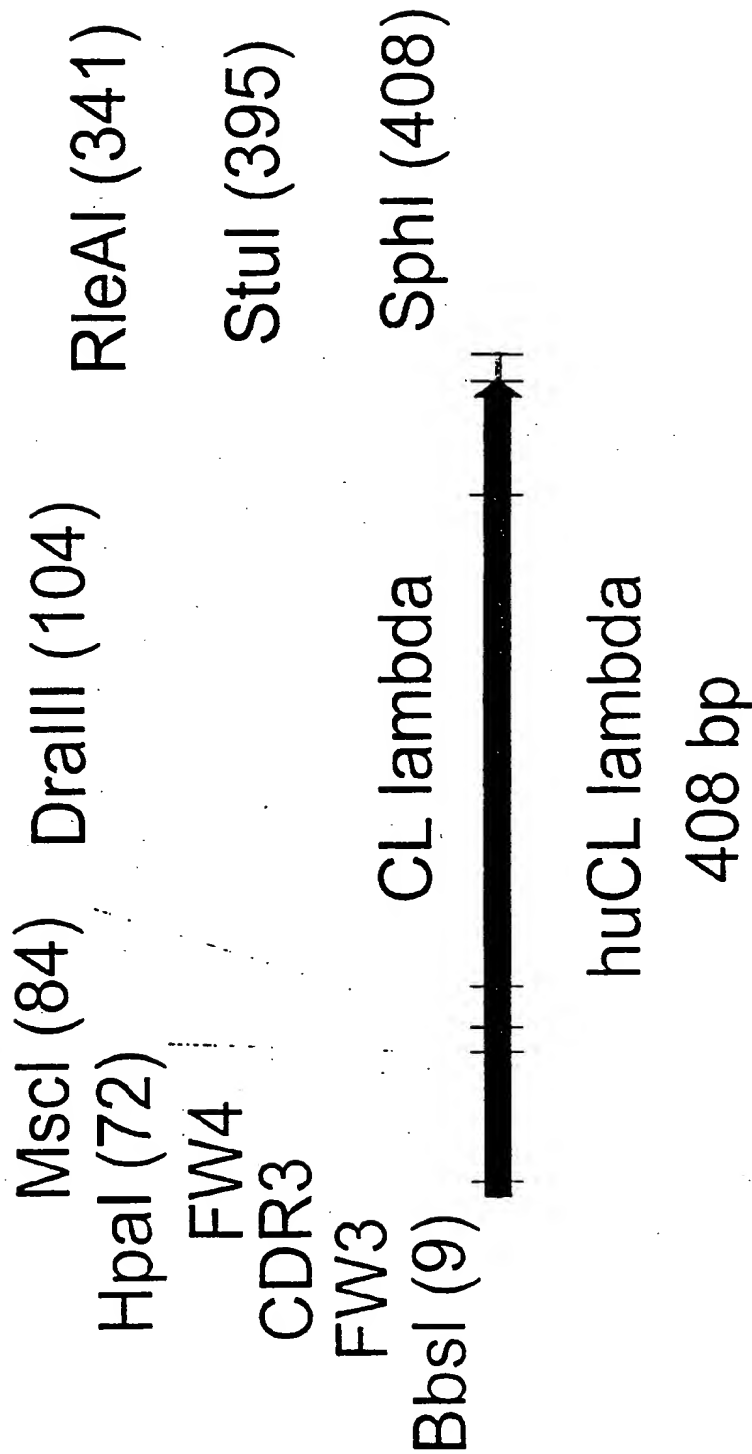


Figure 7C: functional map and sequence of module 24 comprising the synthetic CI gene segment (huCl lambda) (continued)

|     |             |             |            |            |            |
|-----|-------------|-------------|------------|------------|------------|
|     | Bbs I       |             |            |            |            |
|     | ~~~~~       |             |            |            |            |
| 1   | GAAGACGAAG  | CGGATTATTA  | TTGCCAGCAG | CATTATACCA | CCCCGCCTGT |
|     | CTTCTGCTTC  | GCCTAATAAT  | AACGGTCGTC | GTAATATGGT | GGGGCGGACA |
|     |             |             |            |            |            |
|     |             | Hpa I       | Msc I      |            | Dra III    |
|     |             | ~~~~~       | ~~~~~      |            | ~~~~~      |
| 51  | GTTTGGCGGC  | GGCACGAAGT  | TAACCGTTCT | TGGCCAGCCG | AAAGCCGCAC |
|     | CAAACCGCCG  | CCGTGCTTCA  | ATTGGCAAGA | ACCGGTCGGC | TTTCGGCGTG |
|     |             |             |            |            |            |
|     | Dra III     |             |            |            |            |
|     | ~~~~~       |             |            |            |            |
| 101 | CGAGTGTGAC  | GCTGTTTCCG  | CCGAGCAGCG | AAGAATTGCA | GGCGAACAAA |
|     | GCTCACACTG  | CGACAAAGGC  | GGCTCGTCGC | TTCTTAACGT | CCGCTTGTTT |
|     |             |             |            |            |            |
| 151 | GGGACCCCTGG | TGTGCCCTGAT | TAGCGACTTT | TATCCGGGAG | CCGTGACAGT |
|     | CGCTGGGACC  | ACACGGACTA  | ATCGCTGAAA | ATAGGCCCTC | GGCACTGTCA |
|     |             |             |            |            |            |
| 201 | GGCCTGGAAG  | GCAGATAGCA  | GCCCCGTCAA | GGCGGGAGTG | GAGACCACCA |
|     | CCGGACCTTC  | CGTCTATCGT  | CGGGGCAGTT | CCGCCCTCAC | CTCTGGTGGT |

Figure 7C: functional map and sequence of module 24 comprising the synthetic Cl gene segment (huCl lambda) (continued)

251 CACCCCTCCAA ACAAGCAAC AACAGTACG CGGCCAGCAG CTATCTGAGC  
GTGGGAGGTT TGTTTCGTTG TTGTTTCATGC GCCGGTCGTC GATAGACTCG

RleAI

~~~~~

301 CTGACGCCCTG AGCAGTGGAA GTCCACACAGA AGCTACAGCT GCCAGGTCAC  
GACTGCGGGAC TCGTCACCTT CAGGGTGTCT TCGATGTCGA CGGTCCAGTG

StuI

~~~~~

351 GCATGAGGGG AGCACCGTGG AAAAAACCGT TCGGCCGACT GAGGCCCTGAT  
CGTACTCCCC TCGTGGCACC TTTTTTGGCA ACGCGGCTGA CTCCGGACTA

SphI

~~~~~

401 AAGCATGC  
TTCGTACG

Figure 7D: oligonucleotides used for synthesis of module M24 containing Cλ gene segment

M24: assembly PCR

M24-A: GAAGACAAGCGGATTATTATGCCAGCAGCATTATACACCCGCCCTGTGTTGGCGCG-  
GCACGAAGTTAACCGTTC

M24-B: CAATTCTCGCTCGGCGGAACAGCGTCACACTCGGTGGGCTTCGGCTGGCCAA-  
GAACGGTAACTTCGTGCCGC

M24-C: CGCCGAGCAGCGAAGAATTGCAGGGCAACAAGCGACCTGGTGTGCTGATTAGCGACT-  
TTTATCCGGAGCCGTGACA

M24-D: TGTTGGAGGGGTGGTCTCCACITCCCGCCTGACGGGGCTGCTATCGCCTCCAG-  
GCCACTGICACGGCTCCCCG

M24-E: CCACACCCCTCCAAACAAGCAACAAGTACGCGGCCAGCAGCTATCTGAGCCTGACGC-  
CTGAGCAGTGGAAAGTCCCACAGAAGCTACAGCTG

M24-F: GCAITGCTTATCAGGCCCTCAGTCGGCGCAACGGTTTTTCCACGGTGCICCCCCCAIGCGT-  
GACCTGGCAGCTGAGCTTC

Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2

|                                                        |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
|--------------------------------------------------------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|-------|
| M                                                      | K | Q | S | T | I | A | L | A | L | L | P | L | L | F | T | P     |
|                                                        |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | SapI  |
| ~~~~~                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| ATGAAACAAA GCACTATTGC ACTGGCACTC TTACCGTTGC TCTTCACCCC |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| TACTTTGTTT CGTGATAACG TGACCGTGAG AATGGCAACG AGAAGTGGGG |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| ~~~~~                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| V                                                      | T | K | A | D | Y | K | D | E | V | Q | L | V | E | S | G |       |
|                                                        |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | MfeI  |
| ~~~~~                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| TGTTACCAA GCGACTACA AAGATGAAGT GCAATTGGTG GAAAGCGGCG   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| ACAATGGTTT CGGCTGATGT TTCTACTTCA CGTTAACAC CTTTCGCCCG  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| ~~~~~                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| G                                                      | G | L | V | Q | P | G | G | S | L | R | L | S | C | A | A | S     |
|                                                        |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | BspEI |
| ~~~~~                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| GCGGCCTGGT GCAACCGGGC GGCAGCCTGC GTCTGAGCTG CGCGGCCTCC |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| CGCCGGACCA CGTTGGCCCC CCGTCGGACG CAGACTCGAC GCGCCGGAGG |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| ~~~~~                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| G                                                      | F | T | F | S | S | Y | A | M | S | W | V | R | Q | A | P | G     |
|                                                        |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   | BstXI |
| ~~~~~                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| GGATTACCT TTAGCAGCTA TGCATGAGC TGGGTGCGCC AAGCCCCCTGG  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| ~~~~~                                                  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |
| CCTAAATGGA AATCGTCGAT ACGCTACTCG ACCACGCGG TTCGGGGACC  |   |   |   |   |   |   |   |   |   |   |   |   |   |   |   |       |

Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2 (continued)

|                                                          |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
|----------------------------------------------------------|---|---|---|---|---|---|---|---|---|---|---|---|--------|-------|
| K                                                        | G | L | E | W | V | S | A | I | S | G | S | G | S      | T     |
| XhoI                                                     |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
| -----                                                    |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
| GAAGGCTCTC GAGTGGGTGA GCGCGATTAG CCGTAGCGGC GGCAGCACCT   |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
| CTTCCCAGAG CTCACCCACT CCGGCTAATC GCCATCGCCG CCGTCGTGGA   |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
| Y                                                        | Y | A | D | S | V | K | G | R | F | T | I | S | R      | D N S |
|                                                          |   |   |   |   |   |   |   |   |   |   |   |   | PmlI   | NspV  |
|                                                          |   |   |   |   |   |   |   |   |   |   |   |   | -----  | ----  |
| ATTATGCGGA TAGCGTGAAA GGCCGTTTTC CCATTTCACG TGATAATTTCG  |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
| TAAATACGCCCT ATCGCACTTT CCGGCAAAAT GGTAAGTGC ACTATTAAAGC |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
| K                                                        | N | T | L | Y | L | Q | M | N | S | L | R | A | E      | D T A |
|                                                          |   |   |   |   |   |   |   |   |   |   |   |   | NspV   | EagI  |
|                                                          |   |   |   |   |   |   |   |   |   |   |   |   | ----   | ----  |
| AAAACACCC TGTATCTGCA AATGAACAGC CTGCGTGCGG AAGATACGGC    |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
| TTTTTGTGGG ACATAGACGT TTAAGTGTTCG GACGCACGCC TTCTATGCCG  |   |   |   |   |   |   |   |   |   |   |   |   |        |       |
| V                                                        | Y | Y | C | A | R | W | G | G | D | G | F | Y | A      | M D   |
|                                                          |   |   |   |   |   |   |   |   |   |   |   |   | BssHII |       |
|                                                          |   |   |   |   |   |   |   |   |   |   |   |   | -----  |       |
| CGTGTATTAT TCGCGCGGTT GGGCGGCCGA TGGCTTTTAT GCGATGGATT   |   |   |   |   |   |   |   |   |   |   |   |   |        |       |



Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2 (continued)

```

GCACATAATA ACGCGCGCAA CCGCGCGCT ACCGAAAATA CGCTACCTAA
Y W G Q G T L V T V S S A G G G S
 StyI

ATTGGGGCCA AGGCACCCCTG GTGACGGTTA GCTCAGCGGG TGGCGGTTCT
TAACCCCGGT TCCGTGGGAC CACTGCCAAT CGAGTCGCC ACCGCCAAGA
 BlnI

G G G G S G G G G G G G S D I
EcoRV

GGCGGCGGTG GGAGCGGTGG CGGTGGTTCT GCGGTGGTG GTTCCGATAT
CCGCGGCCAC CCTCGCCACC GCCACCAAGA CCGCCACCAC CAAGGCTATA
 EcoRV

V M T Q S P L S L P V T P G E P
 BanII

CGTGATGACC CAGAGCCACC TGAGCCTGCC AGTGA CTCCG GCGAGCCTG
GCACTACTGG GTCTCGGGTG ACTCGGACGG TCACTGAGGC CCGCTCGGAC
 PstI

A S I S C R S S Q S L L H S N G Y
 PstI

CGAGCATTAG CTGCAGAAGC AGCCAAAGCC TGCTGCATAG CAACGGCTAT
GCTCGTAATC GACGCTCTTCG TCGGTTTCGG ACGACGTATC GTTGCCGATA

```

Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2 (continued)

|                                                         |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
|---------------------------------------------------------|---|---|---|---|---|---|---|---|---|----------|---|---|---|---|-------|---|--|
| N                                                       | Y | L | D | W | Y | L | Q | K | P | G        | Q | S | P | Q | L     | L |  |
|                                                         |   |   |   |   |   |   |   |   |   | KpnI     |   |   |   |   | SexAI |   |  |
| -----                                                   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| AAACTATCTGG ATTGGTACCT TCAAAAACCA GTCAAAGCC CGCAGCTATT  |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| TTGATAGACC TAACCATGGA AGTTTGTGTT CCAGTTTCGG GCGTCGATAA  |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| -----                                                   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| I                                                       | Y | L | G | S | N | R | A | S | G | V        | P | D | R | F | S     |   |  |
| AseI                                                    |   |   |   |   |   |   |   |   |   | EcoO109I |   |   |   |   |       |   |  |
| -----                                                   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| AATTATCTG GGCAGCAACC GTGCCAGTGG GGTCCTGGAT CGTTTTCGCG   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| TTAAATAGAC CCGTCGTGG CACGGTCACC CCAGGGCCTA GCAAAATCGC   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| -----                                                   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| G                                                       | S | G | S | G | T | D | F | T | L | K        | I | S | R | V | E     | A |  |
|                                                         |   |   |   |   |   |   |   |   |   | BamHI    |   |   |   |   |       |   |  |
| -----                                                   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| GCTCTGGATC CGGCACCGAT TTTACCTCTGA AAATTAGCCG TGTGGAAGCT |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| CGAGACCTAG GCCGTGGCTA AATGGGACT TTAAATCGGC ACACCTTCGA   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| -----                                                   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| E                                                       | D | V | G | V | Y | Y | C | Q | Q | H        | Y | T | T | P | P     | T |  |
| BbsI                                                    |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| -----                                                   |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| GAAGACGTGG GCGTGTATTA TTGCCAGCAG CATTATACCA CCCC GCCGAC |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |
| CTTCTGCACC CGCACATAAT AACGGTCGTC GTAATATGGT GGGGCGGCTG  |   |   |   |   |   |   |   |   |   |          |   |   |   |   |       |   |  |

Figure 8: sequence and restriction map of the synthetic gene encoding the consensus single-chain fragment VH3-Vk2 (continued)

|             |            |            |             |     |   |   |   |   |   |   |   |   |   |
|-------------|------------|------------|-------------|-----|---|---|---|---|---|---|---|---|---|
| F           | G          | Q          | G           | T   | K | V | E | I | K | R | T | E | F |
| MSCI        |            |            |             |     |   |   |   |   |   |   |   |   |   |
| -----       |            |            |             |     |   |   |   |   |   |   |   |   |   |
| BsiWI ECORI |            |            |             |     |   |   |   |   |   |   |   |   |   |
| -----       |            |            |             |     |   |   |   |   |   |   |   |   |   |
| CTTTGGCCAG  | GGTACGAAAG | TTGAAATTAA | ACGTACGGAA  | TTC |   |   |   |   |   |   |   |   |   |
| GAAACCGGTC  | CCATGCTTC  | AACTTTAATT | TGCATGCCCTT | AAG |   |   |   |   |   |   |   |   |   |

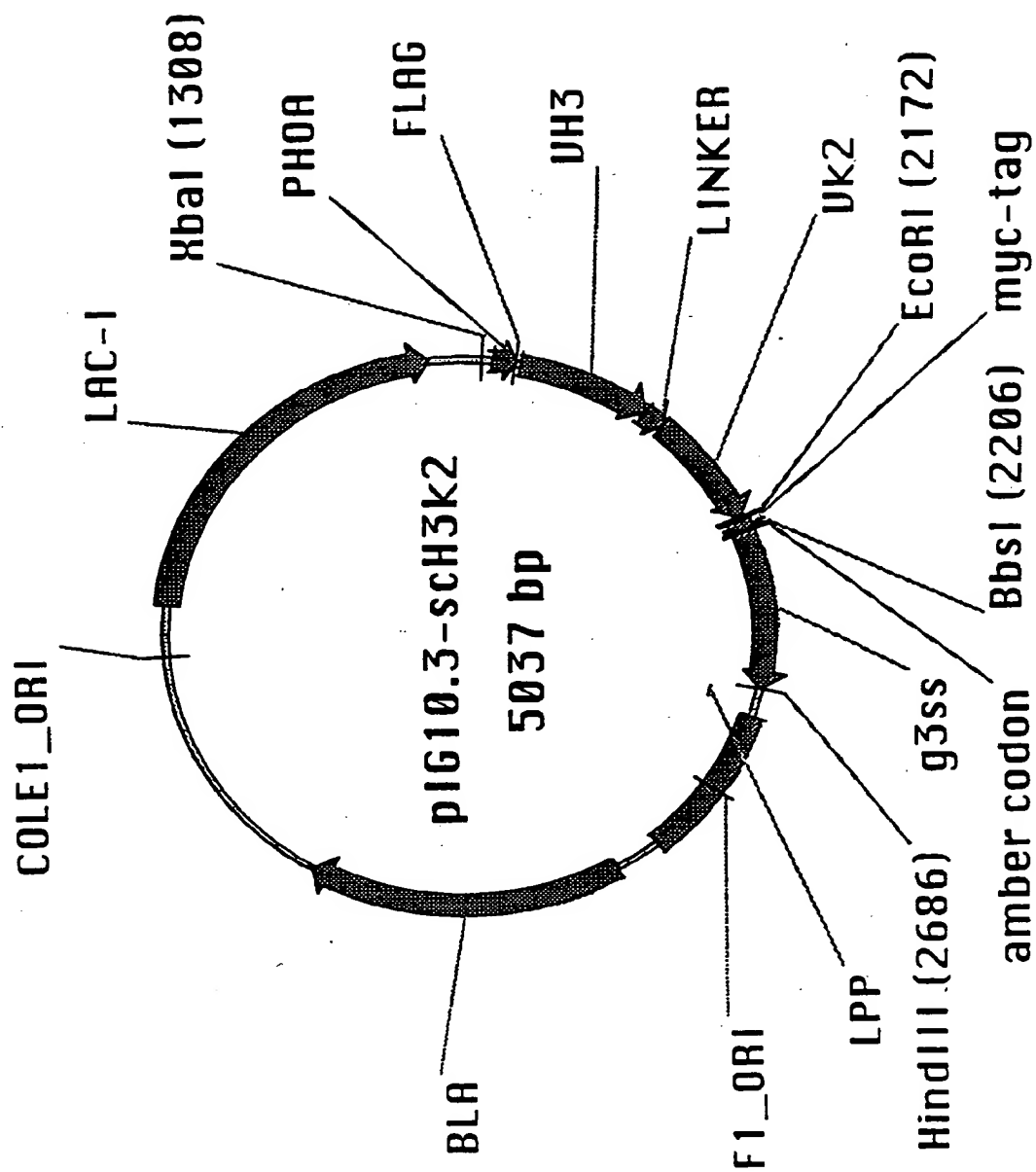


Figure 9: Phage display vector pIG10.3

00420" 49006460

Figure 10: Sequence analysis of initial libraries

|   |      |   |   |   |   |   |   |   |   |   |   |   |   |
|---|------|---|---|---|---|---|---|---|---|---|---|---|---|
| A | 100A | Y | - | - | - | - | - | - | - | - | - | - | - |
|   | 100B | A | - | - | - | - | - | - | - | - | - | - | - |
|   | 100C | - | - | - | - | - | - | - | - | - | - | - | - |
|   | 100D | - | - | - | - | - | - | - | - | - | - | - | - |
|   | 100E | M | - | - | - | - | - | - | - | - | - | - | - |
|   | 101  | D | D | D | D | D | D | D | D | D | D | D | D |
|   | 102  | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y |
|   | 103  | W | W | W | W | W | W | W | W | W | W | W | W |
|   | 99   | G | N | W | Y | A | G | Q | R | N | S | K | A |
|   | 98   | D | M | E | L | K | I | A | T | R | D | F | Q |
|   | 97   | G | K | T | E | L | T | E | I | N | G | T | P |
|   | 96   | G | G | R | R | F | N | N | A | Y | V | K | A |
| B | 95   | W | F | H | V | K | W | I | T | W | S | S | V |
|   | 94   | R | R | R | R | R | R | R | R | R | R | R | R |
|   | 93   | A | A | A | A | A | A | A | A | A | A | A | A |
|   | 92   | C | C | C | C | C | C | C | C | C | C | C | C |
|   | 100  | F | Y | H | H | R | Y | P | - | S | K | A | D |
|   | 100A | Y | - | - | - | - | - | - | - | - | - | - | - |
|   | 100B | A | - | - | - | - | - | - | - | - | - | - | - |
|   | 100C | - | - | - | - | - | - | - | - | - | - | - | - |
|   | 100D | - | - | - | - | - | - | - | - | - | - | - | - |
|   | 100E | M | - | - | - | - | - | - | - | - | - | - | - |
|   | 101  | D | D | D | D | D | D | D | D | D | D | D | D |
|   | 102  | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y | Y |
|   | 103  | W | W | W | W | W | W | W | W | W | W | W | W |

Figure 10: Sequence analysis of initial libraries

|   |   |   |   |   |   |   |   |   |   |   |   |   |   |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| C | C | C | C | C | C | C | C | C | C | C | C | C | C |
| A | A | A | A | A | A | A | A | A | A | A | A | A | A |
| R | R | R | R | R | R | R | R | R | R | R | R | R | R |
| Y | M | K | T | Y | * | R | M | K | S | Y |   |   |   |
| F | A | N | Q | P | G | N | K | G | W | A |   |   |   |
| V | L | Q | S | Y | S | P | P | S | T | G |   |   |   |
| H | R | M | F | R | G | W | M | E | N | T |   |   |   |
| F | A | V | W | S | S | N | L | F | D | T |   |   |   |
| L | S | F | E | N | E | V | N | L | K | F |   |   |   |
| Y | G | H | Q | F | H | N | R | E | P | K |   |   |   |
| T | K | A | Q | F | W | Y | D | T | N | Q |   |   |   |
| M | Y | R | K | M | S | L | G | D | F | G |   |   |   |
| V | I | K | V | P | I | H | T | V | I | P |   |   |   |
| M | M | F | M | M | F | F | M | M | M | M |   |   |   |
| D | D | D | D | D | D | D | D | D | D | D |   |   |   |
| V | V | V | Y | V | V | V | V | Y | V | Y |   |   |   |
| W | W | W | W | W | W | W | W | W | W | W |   |   |   |

Figure 11: Expression analysis of initial library

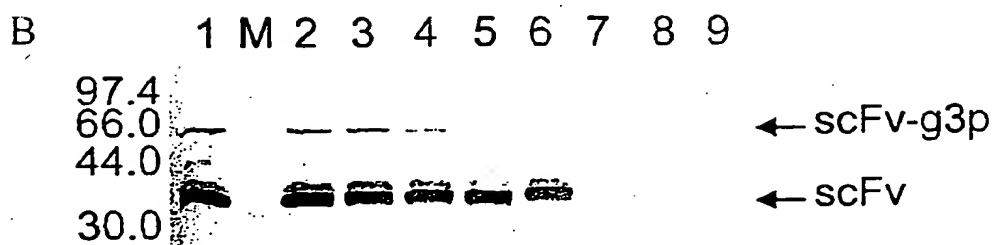
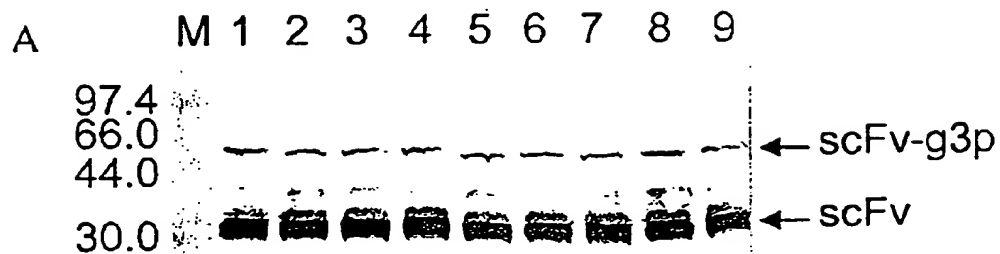


Figure 12: Increase of specificity during the panning rounds

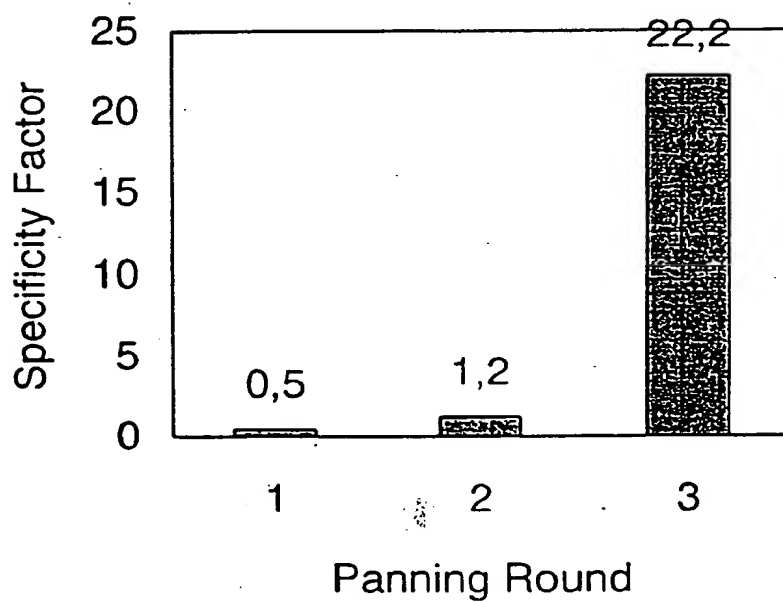




Figure 13: Phage ELISA of clones after the 3rd round of panning

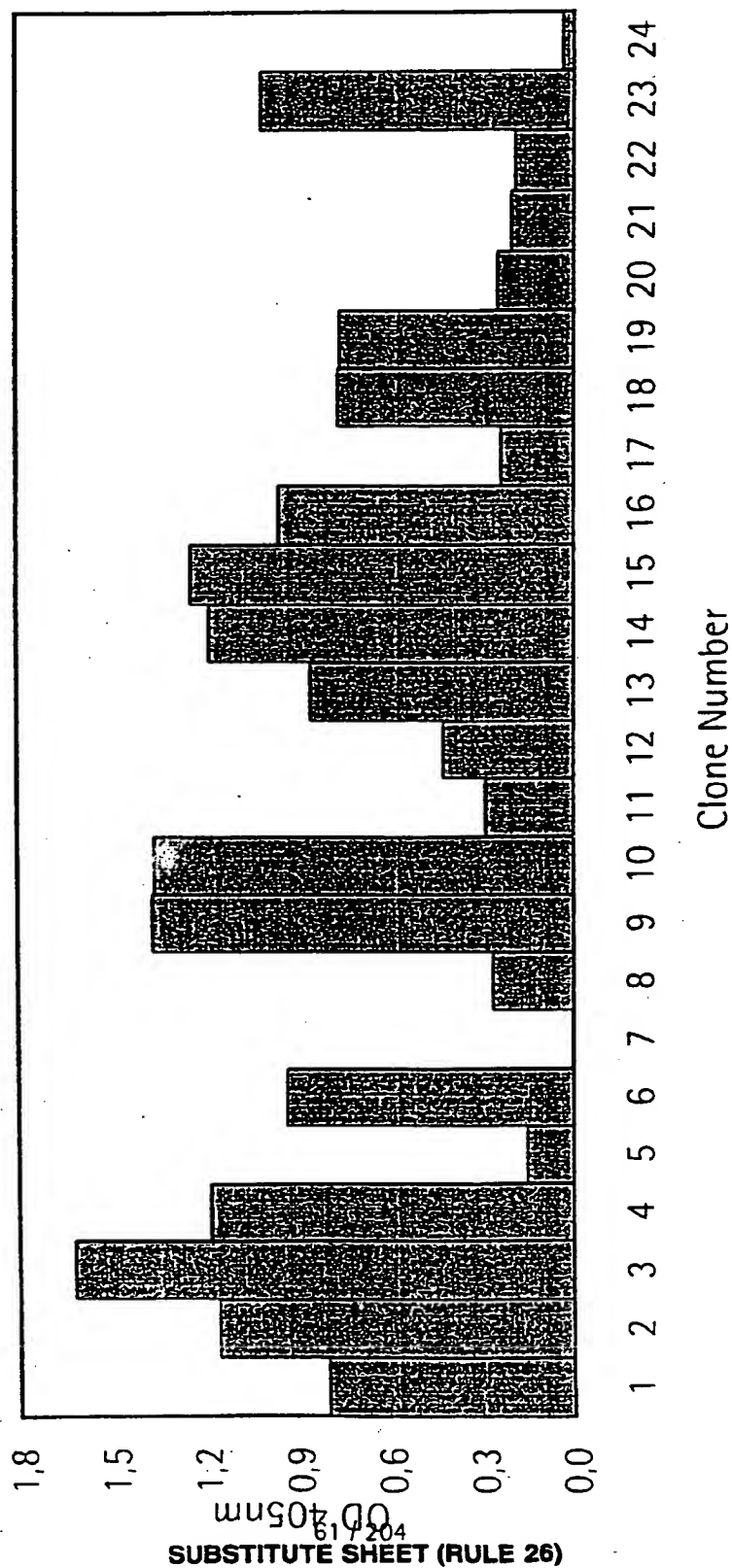
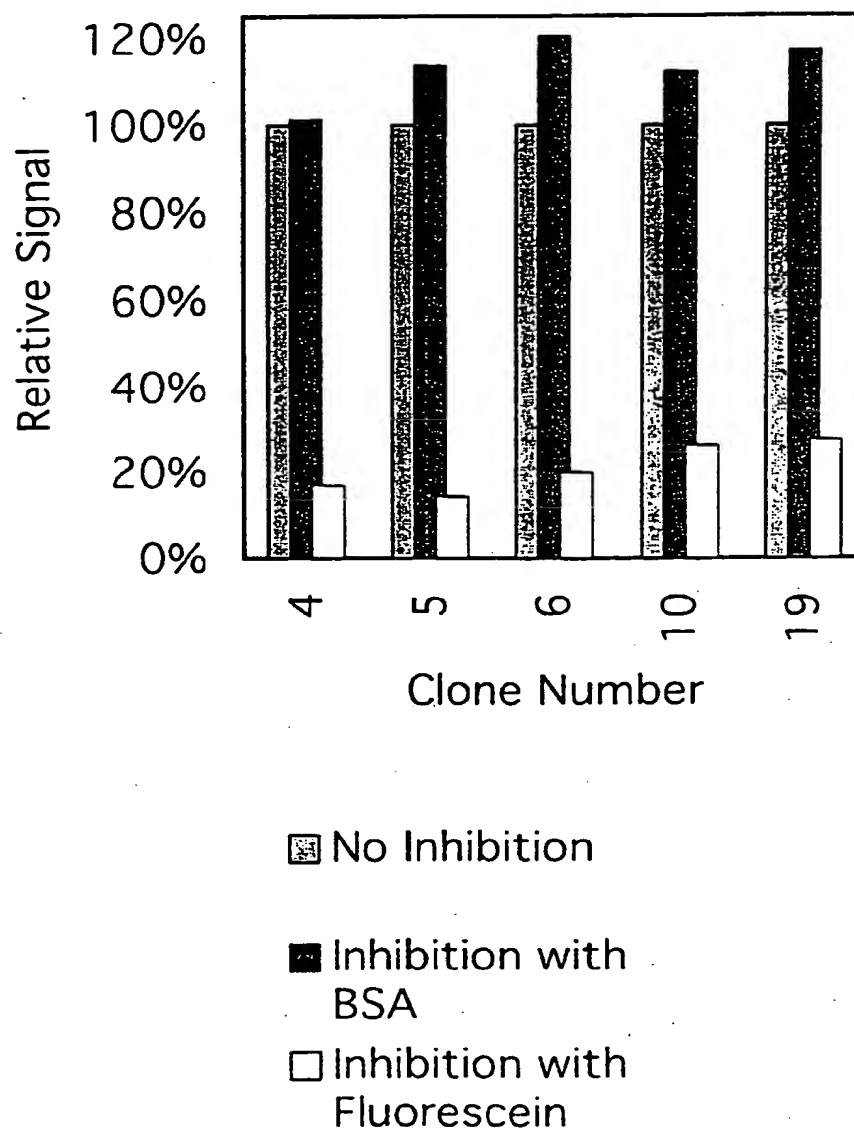


Figure 14: Competition ELISA



| Frequency | 1 | 3 | 1 | 2 | 1 | 1 | 1 | 2 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
|-----------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| 103       | W | W | W | W | W | W | W | W | W | W | W | W | W | W | W | W |
| 102       | V | V | V | V | V | V | V | Y | Y | V | V | V | V | V | V | Y |
| 101       | D | D | D | D | D | D | D | D | D | D | D | D | D | D | D | D |
| 100E      | F | F | F | F | F | M | F | F | F | F | F | F | F | F | F | F |
| 100D      | R | R | R | R | S | Q | V | K | Y | R | R | R | I | Q | R | R |
| 100C      | F | R | H | R | N | D | A | V | K | D | N | P | K | K | A | S |
| 100B      | R | M | R | K | K | F | K | T | V | M | M | R | R | R | F | F |
| 100A      | P | K | L | I | W | S | K | S | R | R | R | A | K | P | S | T |
| 100       | N | R | H | R | K | P | L | Y | S | R | G | F | G | Y | R | Y |
| 99        | Q | K | R | K | M | H | F | R | R | W | R | K | K | T | R | Q |
| 98        | M | Q | K | R | I | V | M | H | M | S | R | K | H | I | K | K |
| 97        | M | K | G | M | K | E | P | F | T | R | P | K | V | H | T | L |
| 96        | R | S | N | K | R | I | K | K | K | K | N | G | M | K | W | K |
| 95        | K | R | R | R | Y | L | R | R | R | K | R | K | R | R | R | K |
| 94        | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R | R |
| 93        | A | A | A | A | A | A | A | A | A | A | A | A | A | A | A | A |
| 92        | C | C | C | C | C | C | C | C | C | C | C | C | C | C | C | C |

Figure 15: Sequence analysis of fluorescein binders

Figure 16: Purification of fluorescein binding scFv fragments

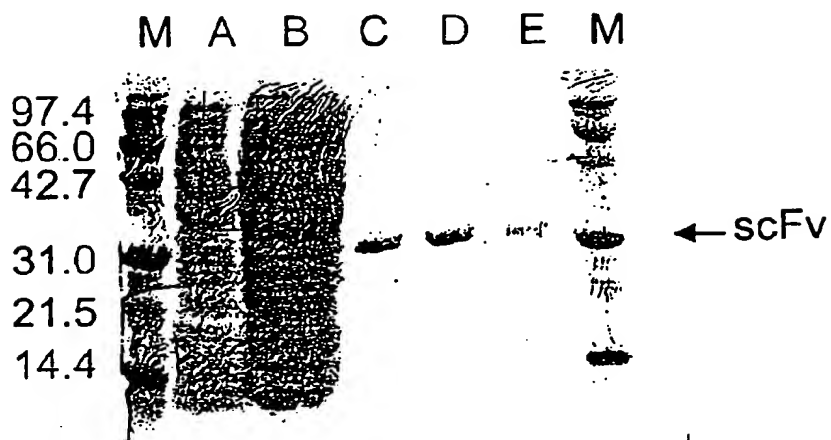


Figure 17: Enrichment factors after three rounds of panning

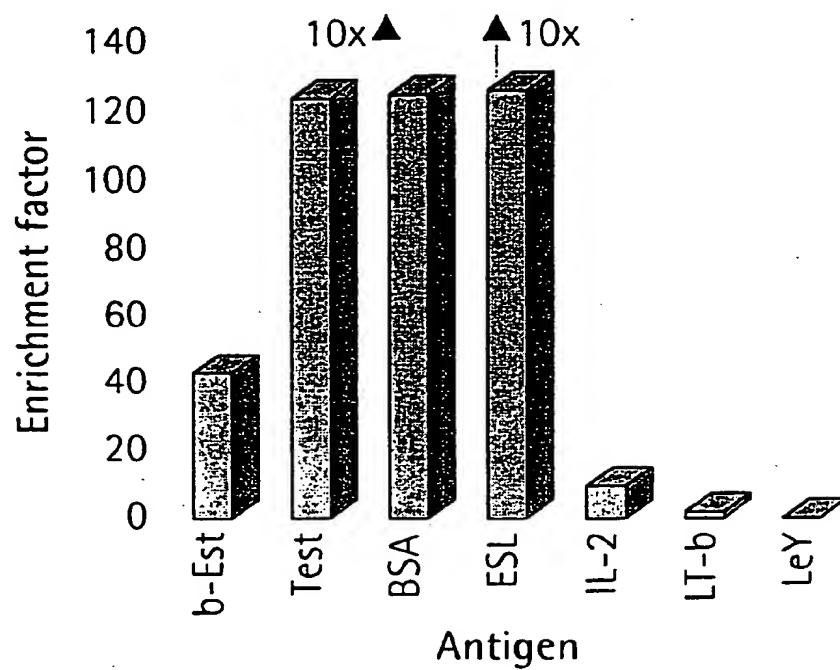


Figure 18: ELISA of anti-ESL-1 and anti- $\beta$ -estradiol antibodies

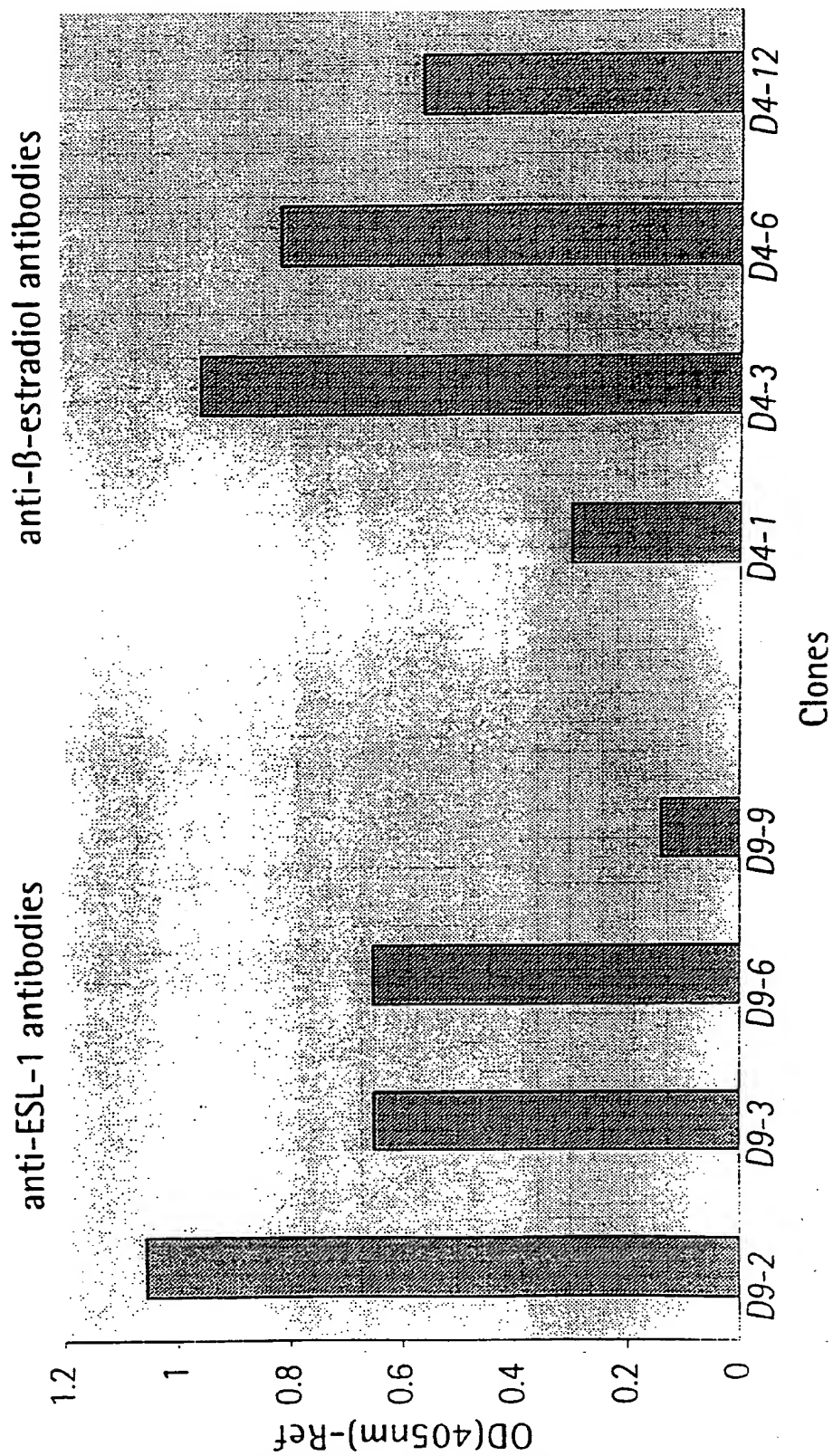
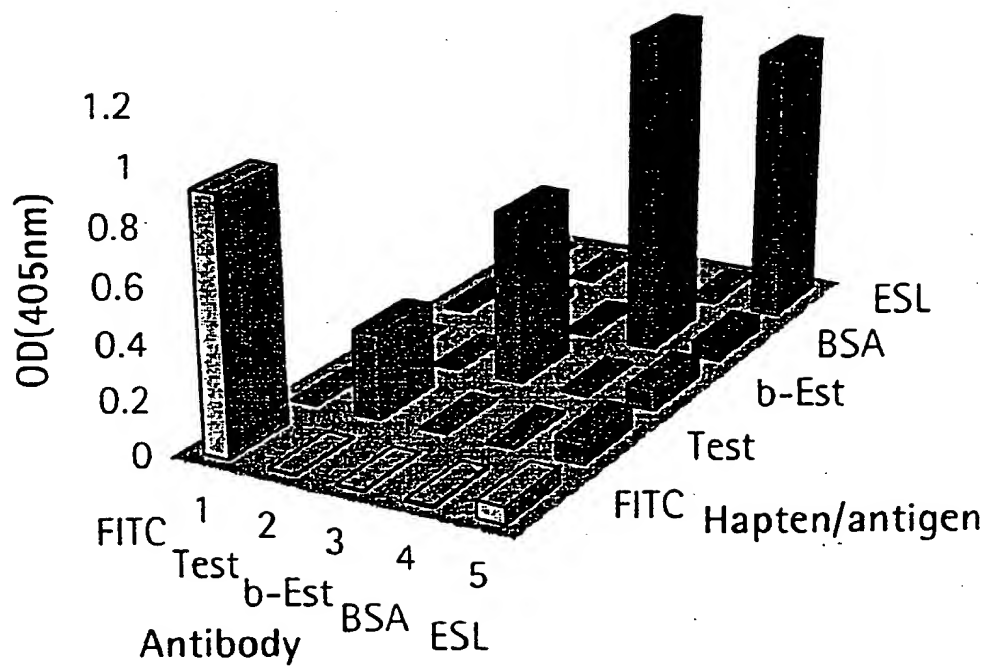


Figure 19: Selectivity and cross-reactivity of HuCAL antibodies



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| Frequency | 4 | 3 | 2 | 1 | 1 | 1 |
|-----------|---|---|---|---|---|---|
| 103       | W | W | W | W | W | W |
| 102       | Y | Y | Y | V | Y | Y |
| 101       | D | D | D | D | D | D |
| 100F      | F | F | F | F | F | F |
| 100D      | A | Q | Q | M | W | Q |
| 100C      | L | M | M | T | K | M |
| 100B      | K | K | K | K | M | Q |
| 100A      | R | Q | N | M | I | R |
| 100       | K | W | R | W | R | S |
| 99        | A | A | A | A | R | A |
| 98        | Q | H | Y | G | L | R |
| 97        | K | R | K | R | P | K |
| 96        | I | N | V | K | K | R |
| 95        | Y | Y | Y | Y | R | Y |
| 94        | R | R | R | R | R | R |
| 93        | A | A | A | A | A | A |
| 92        | C | C | C | C | C | C |

Figure 22: Sequence analysis of lymphotoxin- $\beta$  binders

| Frequency | 103 | 102 | 101 | 100F | 100D | 100C | 100B | 100A | 100 | 99 | 98 | 97 | 96 | 95 | 94 | 93 | 92 |
|-----------|-----|-----|-----|------|------|------|------|------|-----|----|----|----|----|----|----|----|----|
| 16        | W   | V   | D   | F    | H    | G    | K    | I    | K   | S  | R  | Y  | R  | Q  | R  | A  | C  |
| 1         | W   | Y   | D   | M    | P    | D    | Y    | S    | N   | F  | D  | R  | W  | -  | R  | A  | C  |
| 1         | W   | Y   | D   | F    | Q    | V    | W    | Y    | N   | D  | L  | D  | A  | M  | R  | A  | C  |
| 1         | W   | Y   | D   | M    | W    | H    | H    | P    | K   | L  | Y  | A  | Q  | L  | R  | A  | C  |
| 1         | W   | Y   | D   | M    | V    | H    | D    | R    | A   | Q  | E  | I  | L  | R  | R  | A  | C  |
| 1         | W   | V   | D   | F    | S    | Q    | T    | F    | Q   | S  | N  | H  | W  | S  | R  | A  | C  |
| 1         | W   | Y   | D   | M    | W    | E    | N    | E    | T   | Q  | F  | H  | D  | V  | R  | A  | C  |
| 1         | W   | Y   | D   | F    | W    | Y    | W    | F    | I   | L  | T  | P  | W  | D  | R  | A  | C  |

[illegible]

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Figure 24: Sequence analysis of BSA binders

| Frequency |   |
|-----------|---|
| 103       | W |
| 102       | Y |
| 101       | D |
| 100E      | M |
| 100D      | V |
| 100C      | Y |
| 100B      | D |
| 100A      | I |
| 100       | A |
| 99        | Y |
| 98        | F |
| 97        | G |
| 96        | Q |
| 95        | D |
| 94        | R |
| 93        | A |
| 92        | C |
|           | C |
|           | C |
|           | C |
|           | C |
|           | C |
|           | C |

Figure 25: modular pCAL vector system

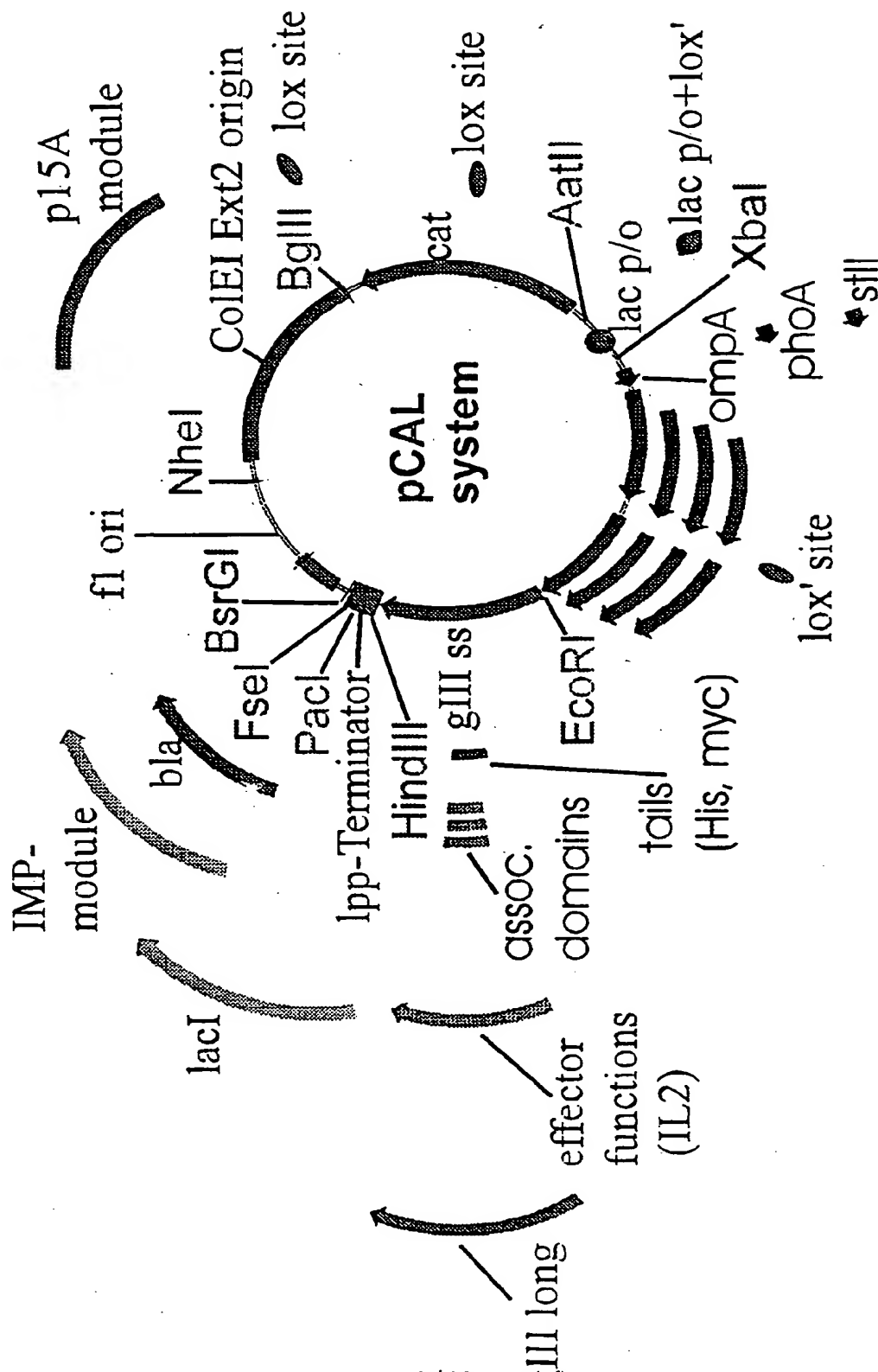


Figure 25a: List of unique restriction sites used in or suitable for HuCAL genes or pCAL vectors

| unique restriction site | Isoschizomers                     |
|-------------------------|-----------------------------------|
| AatII                   | /                                 |
| AflII                   | BfrI, BspTI, Bst98I               |
| AscI                    | /                                 |
| Asel                    | Vspl, AsnI, PshBI                 |
| BamHI                   | BstI                              |
| BbeI                    | EheI, KasI, NarI                  |
| BbsI                    | BpuAI, BpiI                       |
| BglII                   | /                                 |
| BlpI                    | Bpu1102I, CelII, BlpI             |
| BsaBI                   | MamI, Bsh1365I, BsrBRI            |
| BsiWI                   | Pfl23II, SphI, SnaI               |
| BspEI                   | AccIII, BseAI, BsiMI, Kpn2I, MroI |
| BsrGI                   | Bsp1407I, SspBI                   |
| BssHII                  | PaulI                             |
| BstEII                  | BstPI, Eco91I, EcoO65I            |
| BstXI                   | /                                 |
| Bsu36I                  | AocI, CvnI, Eco81I                |
| DraIII                  | /                                 |
| DsmAI                   |                                   |
| EagI                    | BstZI, EclXI, Eco52I, XmaIII      |
| Eco57I                  | /                                 |
| EcoO109I                | DraII                             |
| EcoRI                   | /                                 |
| EcoRV                   | Eco32I                            |
| FseI                    | /                                 |
| HindIII                 | /                                 |
| HpaI                    | /                                 |
| KpnI                    | Acc65I, Asp718I                   |
| MluI                    | /                                 |
| MseI                    | BalI, MluNI                       |

Figure 25a: List of unique restriction sites used in or suitable for HuCAL genes or pCAL vectors

| unique restriction site | Isoschizomers                      |
|-------------------------|------------------------------------|
| MunI                    | MfeI                               |
| NheI                    | /                                  |
| NsiI                    | Ppu10I, EcoT22I, Mph1103I          |
| NspV                    | Bsp119I, BstBI, Csp45I, Lspl, SfuI |
| PacI                    | /                                  |
| PmeI                    | /                                  |
| PmlI                    | BbrPI, Eco72I, PmaCI               |
| Psp5II                  | PpuMI                              |
| PstI                    | /                                  |
| RsrII                   | (RsrI), CpoI, CspI                 |
| SanDI                   | /                                  |
| SapI                    | /                                  |
| SexAI                   | /                                  |
| SpeI                    | /                                  |
| SfiI                    | /                                  |
| SphI                    | BbuI, PaeI, NspI                   |
| StuI                    | AatI, Eco147I                      |
| StyI                    | Eco130I, EcoT14I                   |
| XbaI                    | BspLU11II                          |
| XhoI                    | PaeR7I                             |
| XmaI                    | AvaI, SmaI, Cfr9I, PspAI           |

Figure 26: list of pCAL vector modules

| No   | module/flanking restriction sites | functional element                                            | sites to be removed | sites to be inserted | template      | reference                                                                                                                           |
|------|-----------------------------------|---------------------------------------------------------------|---------------------|----------------------|---------------|-------------------------------------------------------------------------------------------------------------------------------------|
| M1   | AatII-lacp/o-XbaI                 | lac promoter/operator                                         | 2x VspI (AseI)      | AatII                | vector pASK30 | Skerra et al. (1991) Bio/Technology 9, 273-278                                                                                      |
| M2   | BglII-lox-AatII                   | Cre/lox recombination site                                    | 2x VspI (AseI)      | lox, BglII           | (synthetic)   | Hoess et al. (1986) Nucleic Acids Res. 2287-2300                                                                                    |
| M3   | XbaI-lox'-SphI                    | Cre/lox' recombination site                                   | none                | lox', SphI           | (synthetic)   | see M2                                                                                                                              |
| M7-I | EcoRI-glllong-HindIII             | gllp of filamentous phage with N-terminal myctail/amber codon | SphI, BamHI         | none                 | vector pLG10  | Ge et al., (1994) Expressing antibodies in E. coli. In: Antibody engineering: A practical approach. IRL Press, New York, pp 229-266 |



Figure 26: list of pCAL vector modules

|         |                      |                                                                          |                            |                   |              |          |
|---------|----------------------|--------------------------------------------------------------------------|----------------------------|-------------------|--------------|----------|
| M7-II   | EcoRI-gIIIss-HindIII | truncated gIIIp of filamentous phage with N-terminal Gly-Ser linker      | SphI                       |                   | vector pIG10 | see M7-I |
| M7-III  | EcoRI-gIIIss-HindIII | truncated gIIIp of filamentous phage with N-terminal myctail/amber codon | SphI, BbsI                 |                   | vector pIG10 | see M7-I |
| M8      | SphI-lox-HindIII     | Cre/lox recombination site                                               | none                       | lox               | (synthetic)  | see M3   |
| M9-II   | HindIII-lpp-PacI     | lpp-terminator                                                           | none                       | PacI, FseI        | (synthetic)  | see M1   |
| M10-II  | PacI/FseI-bla-BsrGI  | beta-lactamase/bla (ampR)                                                | Vspl, Eco57I, BssSI        | PacI, FseI, BsrGI | pASK30       | see M1   |
| M11-II  | BsrGI-f1 ori-NheI    | origin of single-stranded replication                                    | DrallI (BanII not removed) | BsrGI, NheI       | pASK30       | see M1   |
| M11-III | BsrGI-f1 ori-NheI    | origin of single-stranded replication                                    | DrallI, BanII              | BsrGI, NheI       | pASK30       | see M1   |

[illegible][illegible]

Figure 26: list of pCAL vector modules

|     |                       |                                            |                                                    |  |             |                                                                               |
|-----|-----------------------|--------------------------------------------|----------------------------------------------------|--|-------------|-------------------------------------------------------------------------------|
| M21 | XbaI-stII-SapI        | heat-stable enterotoxin II signal sequence | (synthetic)                                        |  | (synthetic) | Lee et al. (1983) Infect. Immunol. 264-268                                    |
| M41 | AflII-lacI-NheI       | lac-repressor                              | BstXI, MluI, BbsI, BanII, BstEII, HpaI, BbeI, VspI |  | pASK30      | see M1                                                                        |
| M42 | EcoRI-Histail-HindIII | poly-histidine tail                        | (synthetic)                                        |  | (synthetic) | Lindner et al., (1992) Methods: a companion to methods in enzymology 4, 41-56 |

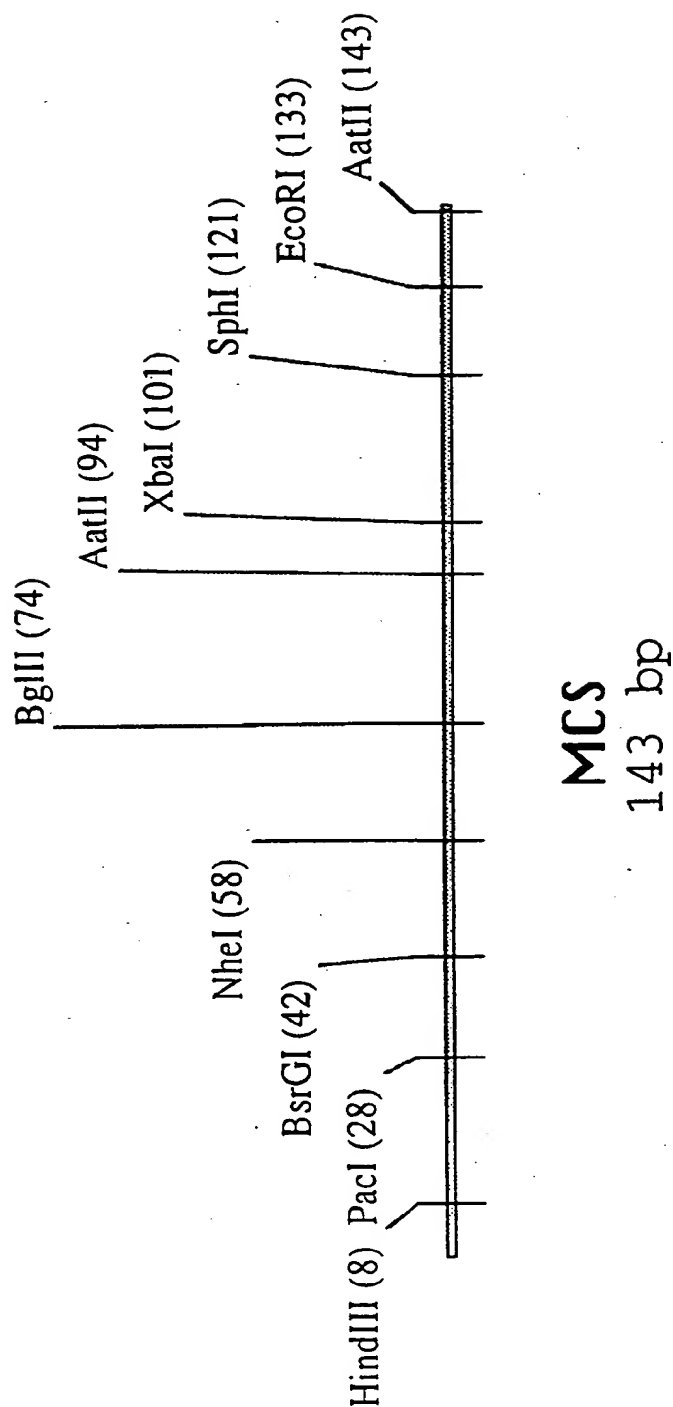


Figure 27: functional map and sequence of MCS module

Figure 27: functional map and sequence of MCS module (continued)

|     | HindIII                                                 | PacI  | BsrGI       |
|-----|---------------------------------------------------------|-------|-------------|
|     | ~~~~~                                                   | ~~~~~ | ~~~~~       |
| 1   | ACATGTAAGC TTCCCCCCCC CCTTAATTAA CCCCCCCCCC TGTACACCCC  |       |             |
|     | TGTACATTTCG AAGGGGGGGG GGAATTAAAT GGGGGGGGGG ACATGTGGGG |       |             |
|     |                                                         |       |             |
|     | NheI                                                    | BglII | AatII XbaI  |
|     | ~~~~~                                                   | ~~~~~ | ~~~~~       |
| 51  | CCCCCGGCTA GCCCCCCCCC CCAGATCTCC CCCCCCCCCG CGTCCCCCCT  |       |             |
|     | GGGGGGCGAT CGGGGGGGGG GGTCTAGAGG GGGGGGGGCT GCAGGGGGGA  |       |             |
|     |                                                         |       |             |
|     | XbaI                                                    | SphI  | EcoRI AatII |
|     | ~~~~~                                                   | ~~~~~ | ~~~~~       |
| 101 | CTAGACCCCC CCCCCGCATG CCCCCCCCCC CGAATTCGAC GTC         |       |             |
|     | GATCTGGGGG GGGGGCGTAC GGGGGGGGGG GCTTAAGCTG CAG         |       |             |

Figure 28: functional map and sequence of pMCS cloning vector

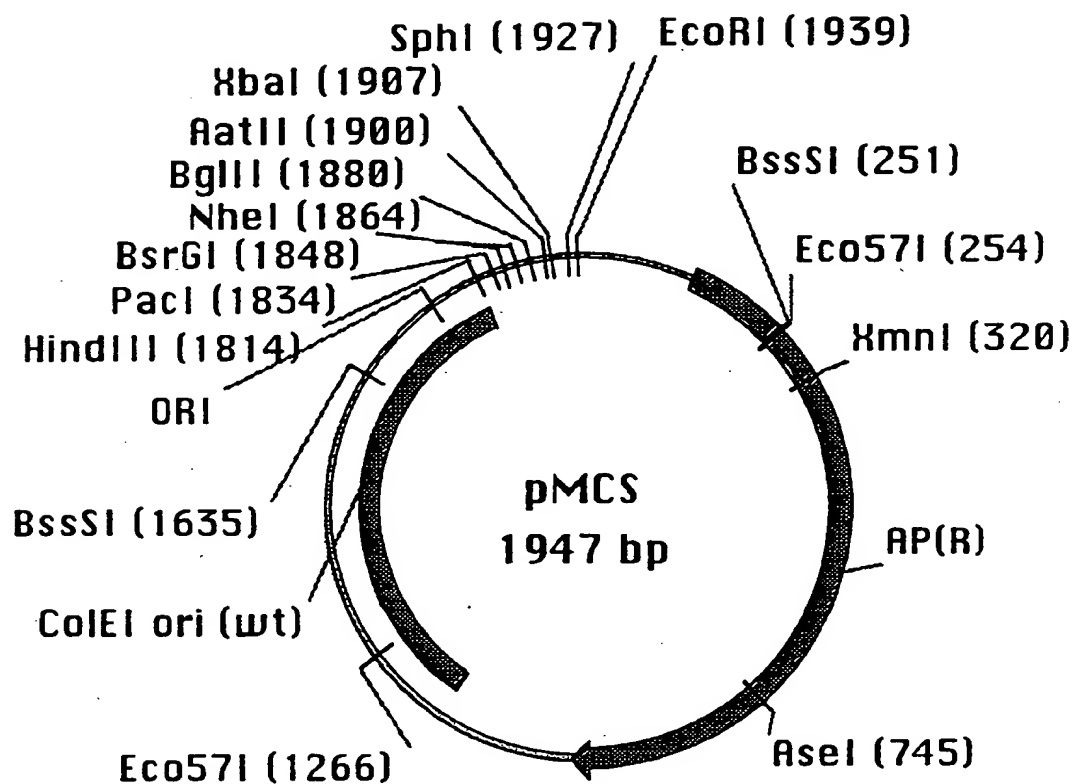


Figure 28: functional map and sequence of pMCS cloning vector (continued)

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1 CAGGTGGCAC TTTTCGGGGA AATGTGCGCG GAACCCCTAT TTGTTTATTT
 GTCCACCCGTG AAAAGCCCCCT TTACACGCGC CTGGGGGATA AACAAATAAA

51 TTCTAAATAC ATTCAAATAT GTATCCGCTC ATGAGACAAT AACCCTGATA
 AAGATTATG TAAGTTTATA CATAGCGGAG TACTCTGTTA TTGGGACTAT

101 AATGCTTCAA TAATATTGAA AAAGGAAGAG TATGAGTATT CAACATTTC
 TTACGAAGTT ATTATAACTT TTTCCTTCTC ATACTCATAA GTTGTAAGG

151 GTGTCGCCCT TATCCCTTT TTTGCGGCAT TTTGCCCTTC TGTTTTGCT
 CACAGCGGGA ATAAGGAAA AACGCCGTA AAACGGAAG ACAAAAACGA

 Eco57I
      ~~~~~

201 CACCCAGAAA CGTGGTGAA AGTAAAGAT GCTGAAGATC AGTTGGGTGC
   GTGGGTCTTT GCGACCACTT TCATTTTCTA CGACTTCTAG TCAACCCACG
      BssSI

251 ACGAGTGGGT TACATCGAAC TGGATCTCAA CAGCGGTAAG ATCCTTGAGA
   TGCTCACCCA ATGTAGCTTG ACCTAGAGTT GTCGCCATTC TAGGAACTCT
      BssSI
      ~~~~~

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Figure 28: functional map and sequence of pMCS cloning vector (continued)

| XmnI  |                                                                                                                    |
|-------|--------------------------------------------------------------------------------------------------------------------|
| ~~~~~ |                                                                                                                    |
| 301   | GTTTTCGCC CGAAGAACGT TTTCCAATGA TGAGCACTTT TAAAGTTCTG<br>CAAAAGCGGG GCTTCTTGCA AAAGTTACT ACTCGTGAAA ATTTCAAGAC     |
| 351   | CTATGTGGCG CGGTATTATC CCGTATTGAC GCCGGGCAAG AGCAACTCGG<br>GATACACCGC GCCATAATAG GGCATAACTG CGGCCCGTTC TCGTTGAGCC   |
| 401   | TCGCCGCATA CACTATTCTC AGAATGACTT GGTGAGTAC TCACCAGTCA<br>AGCGGCGGTAT GTGATAAGAG TCTTACTGAA CCAACTCATG AGTGGTCACT   |
| 451   | CAGAAAAGCA TCTTACGGAT GGCATGACAG TAAGAGAATT ATGCAGTGCT<br>GTCTTTTCGT AGAATGCCCTA CCGTACTGTC ATTCTCTTAA TACGTCACGA  |
| 501   | GCCATAACCA TGAGTGATAA CACTGCGGCC AACTTACTTC TGACAACGAT<br>CGGTATTGGT ACTCACTATT GTGACGCCCG TTGAATGAAG ACTGTTGCTA   |
| 551   | CGGAGGACCG AAGGAGCTAA CCGCTTTTTC GCACAACATG GGGGATCATG<br>GCCCTCCTGGC TTCCCTCGATT GGCGAAAAAA CGTGTTGTAC CCCCTAGTAC |
| 601   | TAACTCGCCT TGATCGTTGG GAACCGGAGC TGAATGAAGC CATAACAAAC<br>ATTGAGCGGA ACTAGCAACC CTTGGCCTCG ACTTACTTCG GTATGGTTTG   |
| 651   | GACGAGCGTG ACACCACGAT GCCTGTAGCA ATGGCAACAA CGTTGCGCAA                                                             |



Figure 28: functional map and sequence of pMCS cloning vector (continued)

|      |            |            |             |             |             |
|------|------------|------------|-------------|-------------|-------------|
|      | CTGCTCGCAC | TGTGGTGCTA | CGGACATCGT  | TACCGTTGTT  | GCAACGCGTT  |
|      |            |            |             |             | AseI        |
|      |            |            |             |             | ~~~~~       |
| 701  | ACTATTAACT | GGCGAACTAC | TTACTCTAGC  | TTCCCCGGCAA | CAATTAATAG  |
|      | TGATAATTGA | CCGCTTGATG | AATGAGATCG  | AAGGGCCGTT  | GTTAATTATC  |
| 751  | ACTGGATGGA | GGCGGATAAA | GTTGCAGGAC  | CACCTTCTGCG | CTCGGCCCTT  |
|      | TGACCTACCT | CCGCCTATTT | CAACGTCCCTG | GTGAAGACGC  | GAGCCGGGAA  |
| 801  | CCGGCTGGCT | GGTTTATTGC | TGATAAATCT  | GGAGCCGGTG  | AGCGTGGGTC  |
|      | GGCCGACCGA | CCAAATAACG | ACTATTTAGA  | CCTCGGCCAC  | TCGCACCCAG  |
| 851  | TCGCGGTATC | ATTGCAGCAC | TGGGGCCAGA  | TGGTAAGCCC  | TCCCCGTATCG |
|      | AGCGCCATAG | TAACGTCGTG | ACCCCGGTCT  | ACCATTCGGG  | AGGGCATAGC  |
| 901  | TAGTTATCTA | CACGACGGGG | AGTCAGGCCAA | CTATGGATGA  | ACGAAATAGA  |
|      | ATCAATAGAT | GTGCTGCCCC | TCAGTCCGTT  | GATACCTACT  | TGCTTTATCT  |
| 951  | CAGATCGCTG | AGATAGGTGC | CTCACTGATT  | AAGCATTGGT  | AACTGTCAGA  |
|      | GTCTAGCGAC | TCTATCCACG | GAGTGACTAA  | TTCGTAACCA  | TTGACAGTCT  |
| 1001 | CCAAGTTTAC | TCATATATAC | TTTAGATTGA  | TTTAAAACCTT | CATTTTAAAT  |
|      | GGTTCAAATG | AGTATATATG | AAATCTAACT  | AAATTTTGAA  | GTAAAAAATTA |

Figure 28: functional map and sequence of pMCS cloning vector (continued)

|      |             |            |             |            |             |
|------|-------------|------------|-------------|------------|-------------|
| 1051 | TTAAAAGGAT  | CTAGGTGAAG | ATCCTTTTGG  | ATAATCTCAT | GACCAAAATC  |
|      | AATTTTCCTA  | GATCCACTTC | TAGGAAAAAC  | TATTAGAGTA | CTGGTTT TAG |
| 1101 | CCTTAACGTG  | AGTTTTCGTT | CCACTGAGCG  | TCAGACCCCG | TAGAAAAGAT  |
|      | GGAAATGCAC  | TCAAAAGCAA | GGTGA CTGC  | AGTCTGGGGC | ATCTTTTCTA  |
| 1151 | CAAAGGATCT  | TCTTGAGATC | CTTTT TTTCT | GCGCGTAATC | TGCTGCTTGC  |
|      | GTTTCCTAGA  | AGAACTCTAG | GAAAAAAGA   | CGCGCATTAG | ACGACGAACG  |
| 1201 | AAACAAAAAA  | ACCACCGCTA | CCAGCGGTGG  | TTTGTTTGCC | GGATCAAGAG  |
|      | TTTGTTT TTT | TGGTGCCGAT | GGTCGCCACC  | AAACAAACGG | CCTAGT TCTC |
| 1251 | CTACCAACTC  | TTTTTCCGAA | GGTAACTGGC  | TTCAGCAGAG | CGCAGATACC  |
|      | GATGGTTGAG  | AAAAAGGCTT | CCATTGACCG  | AAGTCGTCTC | GCGTCTATGG  |
|      |             |            | Eco57I      | ~~~~~      |             |
| 1301 | AAATACTGTC  | CTTCTAGTGT | AGCCG TAGTT | AGGCCACCAC | TTCAAGAACT  |
|      | TTTATGACAG  | GAAGATCACA | TCGGCATCAA  | TCCGGTGGTG | AAGTCTTGA   |
| 1351 | CTGTAGCACC  | GCCTACATAC | CTCGCTCTGC  | TAATCCTGTT | ACCAGTGGCT  |
|      | GACATCGTGG  | CGGATGTATG | GAGCGAGACG  | ATTAGGACAA | TGGTCACCGA  |

Figure 28: functional map and sequence of pMCS cloning vector (continued)

|      |             |             |             |             |             |
|------|-------------|-------------|-------------|-------------|-------------|
| 1401 | GCTGCCAGTG  | CGGATAAGTC  | GTGTCTTACC  | GGGTTGGACT  | CAAGACGATA  |
|      | CGACGGTCAC  | CGCTATTTCAG | CACAGAATGG  | CCCAACCTGA  | GTTCTGCTAT  |
| 1451 | GTTACCGGAT  | AAGCGCAGC   | GGTCGGGCTG  | AACGGGGGGT  | TCGTGCACAC  |
|      | CAATGGCCCTA | TTCCCGCGTCG | CCAGCCCCGAC | TTGCCCCCCCA | AGCACGTGTG  |
| 1501 | AGCCACAGCTT | GGAGCGAACG  | ACCTACACCG  | AAC TGAGATA | CCTACAGCGT  |
|      | TCGGGTCGAA  | CCTCGCTTGC  | TGGATGTGGC  | TTGACTCTAT  | GGATGTGCA   |
| 1551 | GAGCTATGAG  | AAAGCGCCAC  | GCTTCCCAG   | GGGAGAAAGG  | CGGACAGGTA  |
|      | CTCGATACTC  | TTTCGCGGTG  | CGAAGGCTT   | CCCTCTTTCC  | GCCTGTCCAT  |
| 1601 | TCCGGTAAGC  | GGCAGGGTCG  | GAACAGGAGA  | GCGCACGAGG  | GAGCTTCCAG  |
|      | AGGCCATTTCG | CCGTCCCAGC  | CTTGTCCTCT  | CGCGTGCTCC  | CTCGAAGGTC  |
|      |             |             |             | BssSI       |             |
|      |             |             |             | ~~~~~       |             |
| 1651 | GGGGAACGC   | CTGGTATCTT  | TATAGTCCTG  | TCGGGTTTTCG | CCACCTCTGA  |
|      | CCCCTTTGCG  | GACCATAGAA  | ATATCAGGAC  | AGCCCCAAAGC | GGTGGAGACT  |
| 1701 | CTTGAGCGTC  | GATTTTGTG   | ATGCTCGTCA  | GGGGGGCGGA  | GCCTATGGAA  |
|      | GAATCGCAG   | CTAAAAACAC  | TACGAGCAGT  | CCCCCCGCCCT | CGGATACCTT  |
| 1751 | AAACGCCAGC  | AACGGGCCT   | TTTACGGT    | CCTGGCCTT   | TGCTGGCCCTT |

Figure 28: functional map and sequence of pMCS cloning vector (continued)

|      |            |            |             |             |            |
|------|------------|------------|-------------|-------------|------------|
|      | TTTGCGGTCG | TTGCGCCGGA | AAATGCCAA   | GGACCGGAA   | ACGACCGGAA |
|      |            |            |             |             |            |
|      |            | HindIII    |             | PacI        | BsrGI      |
|      |            | ~~~~~      |             | ~~~~~       | ~~~~~      |
| 1801 | TTGCTCACAT | GTAAGCTTCC | CCCCCCCCTT  | AATTAACCCC  | CCCCCCTGTA |
|      | AACGAGTGTA | CATTCGAAGG | GGGGGGGAA   | TTAATTGGGG  | GGGGGACAT  |
|      |            |            |             |             |            |
|      | BsrGI      | NheI       |             | BglII       | AatII      |
|      | ~~         | ~~~~~      |             | ~~~~~       | ~~~~~      |
| 1851 | CACCCCCCCC | CCGCTAGCCC | CCCCCCCCCAG | ATCTCCCCC   | CCCCGACGTC |
|      | GTGGGGGGGG | GGCGATCGGG | GGGGGGGGG   | TAGAGGGGG   | GGGGCTGCAG |
|      |            |            |             |             |            |
|      | XbaI       |            | SphI        |             | EcoRI      |
|      | ~~~~~      |            | ~~~~~       |             | ~~~~~      |
| 1901 | CCCCCTCTAG | ACCCCCCCCC | CGCATGCCCC  | CCCCCCCCGAA | TTCACGT    |
|      | GGGGGAGATC | TGGGGGGGGG | GCGTACGGGG  | GGGGGGGCTT  | AAGTGCA    |



Figure 29: functional map and sequence of pCAL module M1

|     |             |            |                                  |
|-----|-------------|------------|----------------------------------|
|     | AatII       |            |                                  |
|     | ~~~~~       |            |                                  |
| 1   | GACGTCCTTAA | TGTGAGTTAG | CTCACTCATT AGGCACCCCA GGCTTTACAC |
|     | CTGCAGAAATT | ACACTCAATC | GAGTGAGTAA TCCGTGGGGT CCGAAATGTG |
| 51  | TTTATGCTTC  | CGGCTCGTAT | GTTGTGTGGA ATTGTGAGCG GATAACAATT |
|     | AAATACGAAG  | GCCGAGCATA | CAACACACCT TAACACTCGC CTATTGTTAA |
|     |             |            | XbaI                             |
|     |             |            | ~~~~~                            |
| 101 | TCACACAGGA  | AACAGCTATG | ACCATGATTA CGAATTTCTA GA         |
|     | AGTGTGTCCT  | TTGTCGATAC | TGGTACTAAT GCTTAAAGAT CT         |

Figure 30: functional map and sequence of pCAL module M7-II

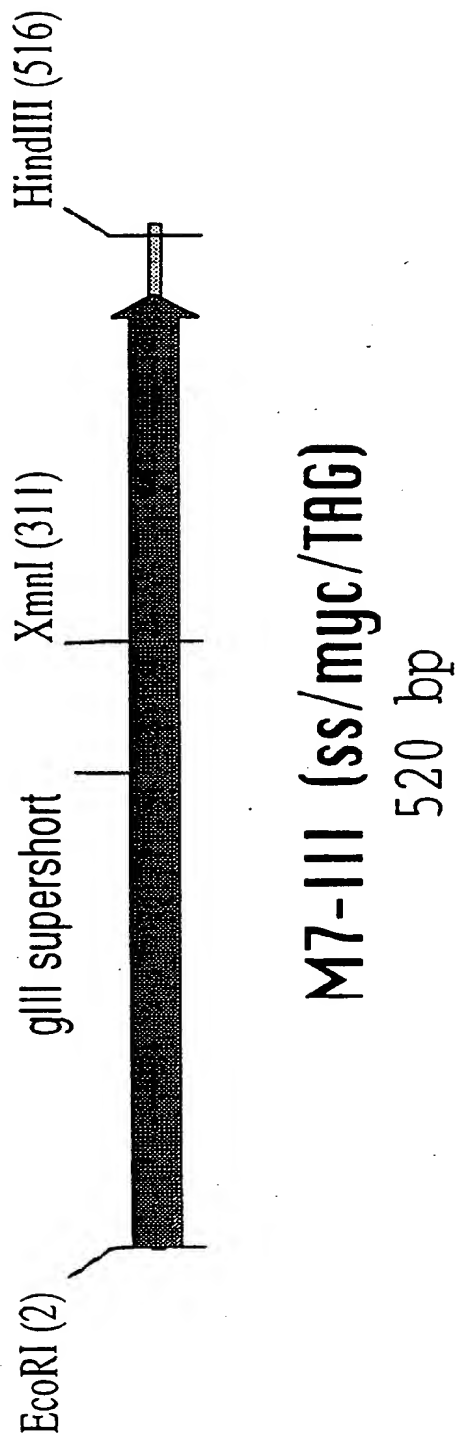


Figure 30: functional map and sequence of pCAL module M7-II (continued)

| EcoRI |                                                         |      |  |
|-------|---------------------------------------------------------|------|--|
|       | ~~~~~                                                   |      |  |
| 1     | GAATTCGAGC AGAAGCTGAT CTCTGAGGAG GATCTGTAGG GTGGTGGCTC  |      |  |
|       | CTTAAGCTCG TCTTCGACTA GAGACTCCTC CTAGACATCC CACCACCGAG  |      |  |
| 51    | TGGTTCGGT GATTTTGATT ATGAAAAGAT GGCAACGCT AATAAGGGGG    |      |  |
|       | ACCAAGGCCA CTAAAACTAA TACTTTTCTA CCGTTTGCGA TTATTCCCCC  |      |  |
| 101   | CTATGACCGA AAATGCCGAT GAAAACGCG TACAGTCTGA CGCTAAAGGC   |      |  |
|       | GATACTGGCT TTTACGGCTA CTTTTCGCG ATGTCAGACT GCGATTTCCG   |      |  |
| 151   | AAACTTGATT CTGTCGCTAC TGATTACGGT GCTGCTATCG ATGGTTTCAT  |      |  |
|       | TTTGAACTAA GACAGCGATG ACTAATGCCA CGACGATAGC TACCAAAAGTA |      |  |
| 201   | TGGTGACGTT TCCGGCCTTG CTAATGGTAA TGGTGCTACT GGTGATTTG   |      |  |
|       | ACCACTGCAA AGGCCGGAAC GATTACCATT ACCACGATGA CCACATAAAC  |      |  |
| 251   | CTGGCTCTAA TTCCCAAATG GCTCAAGTCG GTGACGGTGA TAATCACCT   |      |  |
|       | GACCGAGATT AAGGGTTTAC CGAGTTCAGC CACTGCCACT ATTAAGTGGA  |      |  |
|       |                                                         | XmnI |  |
|       | ~~~~~                                                   |      |  |
| 301   | TTAATGAATA ATTTCCGTCA ATATTACCT TCCCTCCCTC AATCGGTTGA   |      |  |
|       | AATTACTTAT TAAAGGCAGT TATAAATGGA AGGGAGGGAG TTAGCCAACT  |      |  |



Figure 30: functional map and sequence of pCAL module M7-II (continued)

|         |            |            |            |            |            |
|---------|------------|------------|------------|------------|------------|
| 351     | ATGTCGCCCT | TTTGTCTTTG | GCGCTGGTAA | ACCATATGAA | TTTTCTATTG |
|         | TACAGCGGGA | AAACAGAAAC | CGCGACCAAT | TGGTATACTT | AAAAGATAAC |
| 401     | ATTGTGACAA | AATAAACTTA | TTCCGTGGTG | TCTTTGCGTT | TCCTTTATAT |
|         | TAACACTGTT | TTATTGGAAT | AAGGCACCAC | AGAAACGCAA | AGAAATATA  |
| 451     | GTTGCCACCT | TTATGTATGT | ATTTTCTACG | TTTGCTAACA | TACTGCGTAA |
|         | CAACGGTGGA | AATACATACA | TAAAGATGC  | AAACGATTGT | ATGACGCATT |
| HindIII |            |            |            |            |            |
| ~~~~~   |            |            |            |            |            |
| 501     | TAAGGAGTCT | TGATAAGCTT |            |            |            |
|         | ATTCCTCAGA | ACTATTCGAA |            |            |            |

Figure 31: functional map and sequence of pCAL module M9-II

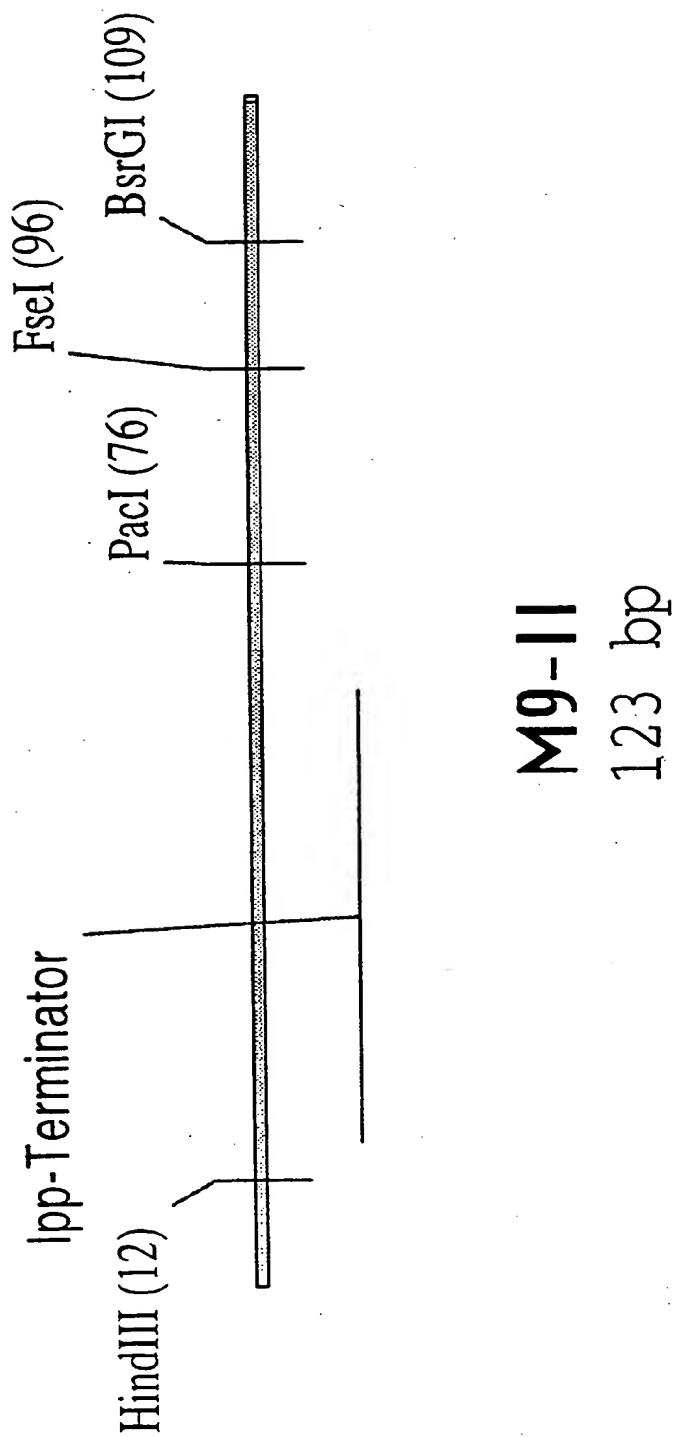


Figure 31: functional map and sequence of pCAL module M9-II (continued)

|     |                                                       |       |       |  |
|-----|-------------------------------------------------------|-------|-------|--|
|     | HindIII                                               |       |       |  |
|     | ~~~~~                                                 |       |       |  |
| 1   | GGGGGGGGG AAGCTGACC TGTGAAGTGA AAAATGGCG AGATTGTGCG   |       |       |  |
|     | CCCCCCCCC TTCGAACTGG AACTTCACT TTTTACCGCG TCTAACACGC  |       |       |  |
|     |                                                       | PacI  | FseI  |  |
|     |                                                       | ~~~~~ | ~~~~~ |  |
| 51  | ACATTTT TGTCTGCCGT TTAATTAAAG GGGGGGGG GCCGGCCTGG     |       |       |  |
|     | TGTAAAAAAA ACAGACGGCA AATTAAATTC CCCCCCCCC CGCCGGGACC |       |       |  |
|     |                                                       |       |       |  |
|     | BsrGI                                                 |       |       |  |
|     | ~~~~~                                                 |       |       |  |
| 101 | GGGGGGGTGT ACAGGGGGGG GGG                             |       |       |  |
|     | CCCCCCCCCA GTCCCCCCCC CCC                             |       |       |  |

Figure 32: functional map and sequence of pCAL module M11-III

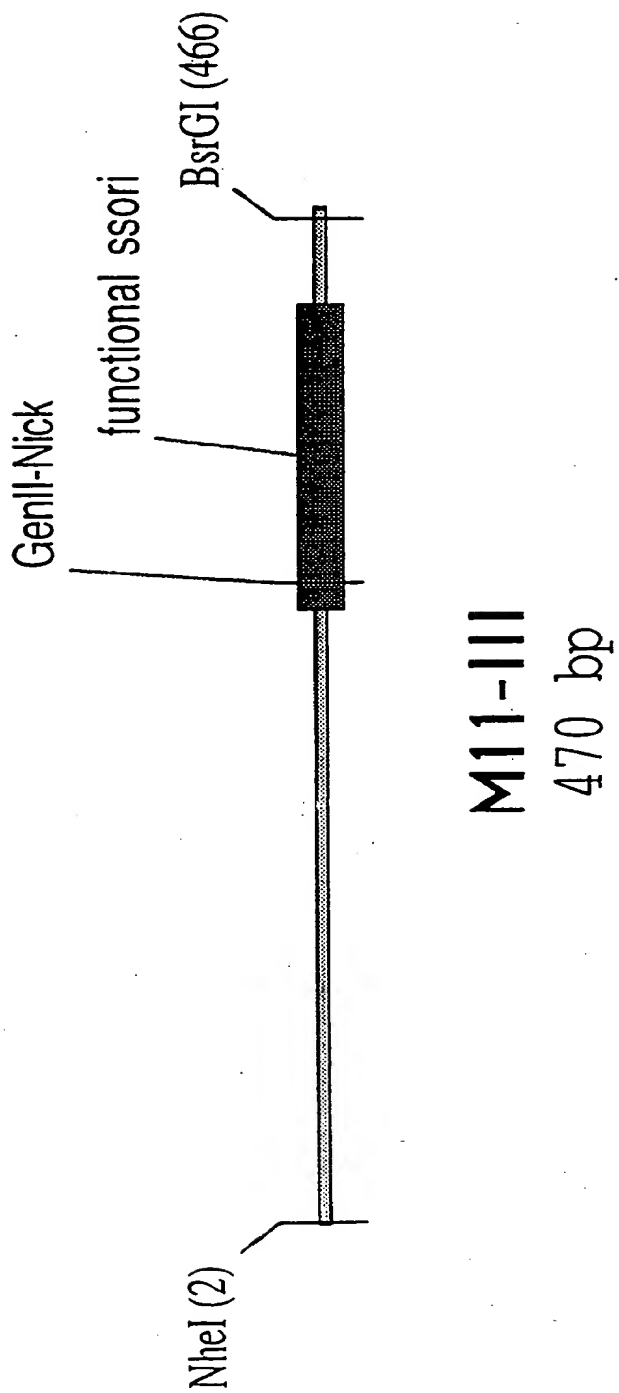


Figure 32: functional map and sequence of pCAL module M11-III (continued)

NheI

~~~~~

|     |            |            |            |             |            |
|-----|------------|------------|------------|-------------|------------|
| 1   | GCTAGCACGC | GCCCTGTAGC | GGCGCATTA  | GGCGGCGGG   | TGTGGTGGTT |
|     | CGATCGTGCG | CGGACATCG  | CCGCGTAATT | CGCGCCGCC   | ACACCACCAA |
| 51  | ACGCGCAGCG | TGACCGCTAC | ACTTGCCAGC | GCCCTAGCG   | CCGCTCCTTT |
|     | TGCGCGTCGC | ACTGGCGATG | TGAACGGTCG | CGGGATCGCG  | GGCGAGGAAA |
| 101 | CGCTTCTTC  | CCTTCCCTTC | TCGCCACGTT | CGCCGGCTTT  | CCCCGTCAAG |
|     | GCGAAGAAG  | GGAAGGAAAG | AGCGGTGCAA | GGCGCCGAAA  | GGGCAGTTC  |
| 151 | CTCTAAATCG | GGGCATCCCT | TTAGGGTCC  | GATTAGTGC   | TTTACGGCAC |
|     | GAGATTAGC  | CCCGTAGGA  | AATCCCAAG  | CTAAATCAG   | AAATGCCGTG |
| 201 | CTCGACCCCA | AAAACTTGA  | TTAGGGTGAT | GGTTCTCGTA  | GTGGGCCATC |
|     | GAGCTGGGGT | TTTTTGAACT | AATCCCACTA | CCAAGAGCAT  | CACCCGGTAG |
| 251 | GCCCTGATAG | ACGGTTTTTC | GCCCTTTGAC | GTTGGAGTCC  | ACGTTCTTTA |
|     | CGGGACTATC | TGCCAAAAG  | CGGAAACTG  | CAACCTCAGG  | TGCAAGAAAT |
| 301 | ATAGTGGAAT | CTTGTTCCTA | ACTGGAACAA | CACCTAACCC  | TATCTCGGTC |
|     | TATCACCTGA | GAACAAGGT  | TGACCTTGTT | GTGAGTTGG   | ATAGAGCCAG |
| 351 | TATTCCTTTG | ATTTATAAGG | GATTTTGCCG | ATTTCCGCCCT | ATTGGTTAAA |

Figure 32: functional map and sequence of pCAL module M11-III (continued)

ATAAGAGAAAC TAAATATTCC CTAAAACGGC TAAAGCCGGA TAACCAATTT

401 AAATGAGCTG ATTTAACAAA AATTTAACGC GAATTTTAAC AAAATATTAA  
TTTACTCGAC TAAATTGTTT TTAAATTGCG CTTAAAATTG TTTTATAATT

BsrGI

~~~~~

451 CGTTTACAAT TTCATGTACA  
GCAAAATGTTA AAGTACATGT

**M14-EXT2**  
**733 bp**

Figure 33: functional map and sequence of pCAL module M14-Ext2 (continued)

BglII  
~~~~~

|     |             |            |             |            |             |
|-----|-------------|------------|-------------|------------|-------------|
| 1   | AGATCTGACC  | AAAATCCCTT | AACGTGAGTT  | TTCGTTCCAC | TGAGCGTCAG  |
|     | CTAGACTGG   | TTTTAGGGAA | TTGCCACTCAA | AAGCAAGGTG | ACTCGCAGTC  |
| 51  | ACCCCGTAGA  | AAAGATCAAA | GGATCTTCTT  | GAGATCCCTT | TTTTCTGCGC  |
|     | TGGGGCATCT  | TTTCTAGTTT | CCTAGAAGAA  | CTCTAGGAAA | AAAAGACGCG  |
| 101 | GTAATCTGCT  | GCTTGCAAAC | AAAAAAACCA  | CCGCTACCAG | CGGTGGTTTG  |
|     | CATTAGACGA  | CGAACGTTTG | TTTTTTTGGT  | GCGATGGTC  | GCCACCAAAC  |
| 151 | TTTGCCGGAT  | CAAGAGCTAC | CAACTCTTTT  | TCCGAAGGTA | ACTGGGTACA  |
|     | AAACGGCCTA  | GTTCTCGATG | GTTGAGAAAA  | AGGCTTCCAT | TGACCGATGT  |
| 201 | GCAGAGCGCA  | GATACCAAAT | ACTGTTCTTC  | TAGTGTAGCC | GTAGTTAGGC  |
|     | CGTCTCGCGT  | CTATGGTTTA | TGACAAGAAG  | ATCACATCGG | CATCAATCCG  |
| 251 | CACCACTTCA  | AGAACTCTGT | AGCACCGCCT  | ACATACCTCG | CTCTGCTAAT  |
|     | GTGGTGAAAGT | TCTTGAGACA | TCGTGGCGGA  | TGTATGGAGC | GAGACGATTA  |
| 301 | CCTGTTACCA  | GTGGCTGCTG | CCAGTGGCGA  | TAAGTCGTGT | CTTACCGGGT  |
|     | GGACAATGGT  | CACCGACGAC | GGTCACCGCT  | ATTCAGCACA | GAATGGCCCCA |
| 351 | TGGACTCAAG  | ACGATAGTTA | CCGATAAGG   | CGCAGCGGTC | GGGCTGAACG  |



Figure 33: functional map and sequence of pCAL module M14-Ext2 (continued)

|     |             |            |             |            |              |
|-----|-------------|------------|-------------|------------|--------------|
|     | ACCTGAGTTC  | TGCTATCAAT | GGCCTATTCC  | GGTCGCCAG  | CCCGACTTGC   |
| 401 | GGGGGTTTCGT | GCACACAGCC | CAGCTTGGAG  | CGAACGACCT | ACACCGAACT   |
|     | CCCCCAAGCA  | CGTGTCGG   | GTCGAACCTC  | GCTTGCTGGA | TGTGGCTTGA   |
| 451 | GAGATACCTA  | CAGCGTGAGC | TATGAGAAAG  | CGCCACGCTT | CCCGAAGGGA   |
|     | CTCTATGGAT  | GTCGCACTCG | ATACTCTTTC  | GCGGTGCGAA | GGGCTTCCCT   |
| 501 | GAAAGCGGA   | CAGTATCCG  | GTAAGCGGCA  | GGTCCGGAAC | AGGAGAGCGC   |
|     | CTTTCCGCCT  | GTCCATAGGC | CATTGCCCCG  | CCCAGCCTTG | TCCTCTCGCG   |
|     |             |            |             | BSSI       | ~            |
| 551 | ACGAGGGAGC  | TTCCAGGGGG | AAACGCCCTGG | TATCTTTATA | GTCCCTGTCCGG |
|     | TGCTCCCCTCG | AAGTCCCCC  | TTTGCGGACC  | ATAGAAATAT | CAGGACAGCC   |
|     | BSSI        |            |             |            |              |
|     | ~~~~~       |            |             |            |              |
| 601 | GTTTCGCCAC  | CTCTGACTTG | AGCGTCGATT  | TTTGTGATGC | TCGTCAGGGG   |
|     | CAAAGCGGTG  | GAGACTGAAC | TCGCAGCTAA  | AAACACTACG | AGCAGTCCCC   |
| 651 | GGCGGAGCCT  | ATGGAAAAC  | GCCAGCAACG  | CGGCCCTTTT | ACGGTTCCCTG  |
|     | CGCCCTCGGA  | TACCTTTTGG | CGGTCGTGTC  | GCCGGAAAAA | TGCCAAGGAC   |

Figure 33: functional map and sequence of pCAL module M14-Ext2 (continued)

701 GCCTTTTGCT GGCCTTTTGC TCACATGGCT AGC  
CGGAAAACGA CCGGAAAACG AGTGTACCGA TCG

NheI  
~~~~~

Figure 34: functional map and sequence of pCAL module M17

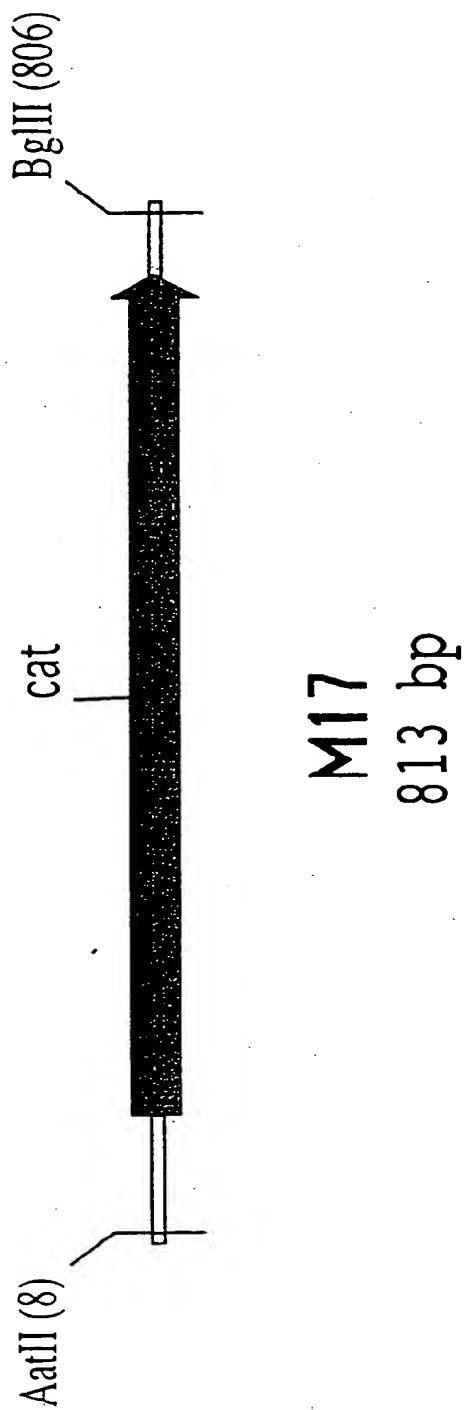


Figure 34: functional map and sequence of pCAL module M17 (continued)

AatII

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1 GGGACGTCGG GTGAGGTTCC AACTTTCACC ATAATGAAAT AAGATCACTA
 CCTGTCAGCC CACTCCAAGG TTGAAAGTGG TATTACTTTA TTCTAGTGAT

51 CCGGGCGGTAT TTTTGTGAGTT ATCGAGATTT TCAGGAGCTA AGGAAGCTAA
 GCGCCGCATA AAAAAGCTCAA TAGCTCTAAA AGTCCCTCGAT TCCTTCGATT

101 AATGGAGAAA AAAATCACTG GATATACCAC CGTTGATATA TCCCAATGGC
 TTACCTCTTT TTTTAGTGAC CTATATGGTG GCAACTATAT AGGTTTACCG

151 ATCGTAAAGA ACATTTTGAG GCATTTTCAGT CAGTTGCTCA ATGTACCTAT
 TAGCATTTCT TGTAAGAACTC CGTAAAGTCA GTCAACGAGT TACATGGATA

201 AACCAGACCG TTCAGCTGGA TATTACGGCC TTTTAAAGA CCGTAAAGAA
 TTGGTCTGGC AAGTCGACCT ATAATGCCGG AAAAATTTCT GGCATTTCTT

251 AAATAAGCAC AAGTTTATC CGCCCTTTAT TCACATTTCT GCCCGCCTGA
 TTTATTCGTG TTCAAAATAG GCCGGAATA AGTGTAAGAA CGGGCGGACT

301 TGAATGCTCA CCCGGAGTTC CGTATGGCAA TGAAAGACGG TGAGCTGGTG
 ACTTACGAGT GGGCCTCAAG GCATACCGTT ACTTCTGCC ACTCGACCCAC

351 ATATGGGATA GTGTTCACCC TTGTTACACC GTTTTCCATG AGCAAACTGA

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Figure 34: functional map and sequence of pCAL module M17 (continued)

|     |             |            |            |             |             |
|-----|-------------|------------|------------|-------------|-------------|
|     | TATACCCCTAT | CACAAGTGGG | AACAATGTGG | CAAAAGGTAC  | TCGTTTGACT  |
| 401 | AACGTTTTCA  | TCGCTCTGGA | GTGAATACCA | CGACGATTTC  | CGGCAGTTTC  |
|     | TTGCAAAAAGT | AGCGAGACCT | CACTTATGGT | GCTGCTAAAG  | GCCGTCAAAG  |
| 451 | TACACATATA  | TTCGCAAGAT | GTGGCGTGT  | ACGGTGAAA   | CCTGGCCCTAT |
|     | ATGTGTATAT  | AAGCGTTCTA | CACCGCACAA | TGCCACTTTT  | GGACCGGATA  |
| 501 | TTCCCTAAAG  | GGTTTATTGA | GAATATGTTT | TTCGTCTCAG  | CCAATCCCCTG |
|     | AAGGATTTC   | CCAAATAACT | CTTATACAAA | AAGCAGAGTC  | GGTTAGGGAC  |
| 551 | GGTGAGTTTC  | ACCAGTTTGG | ATTTAAACGT | AGCCAATATG  | GACAACTTCT  |
|     | CCACTCAAAG  | TGGTCAAAAC | TAAATTTGCA | TCGGTTATAC  | CTGTTGAAGA  |
| 601 | TCGCCCCCGT  | TTTCACTATG | GGCAAATATT | ATACGCAAGG  | CGACAAGGTG  |
|     | AGCGGGGGCA  | AAAGTGATAC | CCGTTTATAA | TATGCGTTCC  | GCTGTTCCAC  |
| 651 | CTGATGCCGC  | TGGCGATTCA | GGTTCATCAT | GCCGTTTGTG  | ATGGCTTCCA  |
|     | GACTACGGCG  | ACCGCTAAGT | CCAAGTAGTA | CGGCAAAACAC | TACCGAAGGT  |
| 701 | TGTCGGCAGA  | ATGCTTAATG | AATTACAACA | GTACTGCCAT  | GAGTGGCAGG  |
|     | ACAGCCGTCT  | TACGAATTAC | TTAATGTTGT | CATGACGCTA  | CTCACCGTCC  |
| 751 | GCGGGGCGTA  | ATTTTTTTAA | GGCAGTTATT | GGGTGCCCTT  | AAACGCCCTG  |

00420"49006760

Figure 34: functional map and sequence of pCAL module M17 (continued)

CGCCCCCGCAT TAAAAAATT CCGTCAATAA CCCACGGGAA TTTGCGGACC

BglII

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801 TGCTAGATCT TCC  
ACGATCTAGA AGG

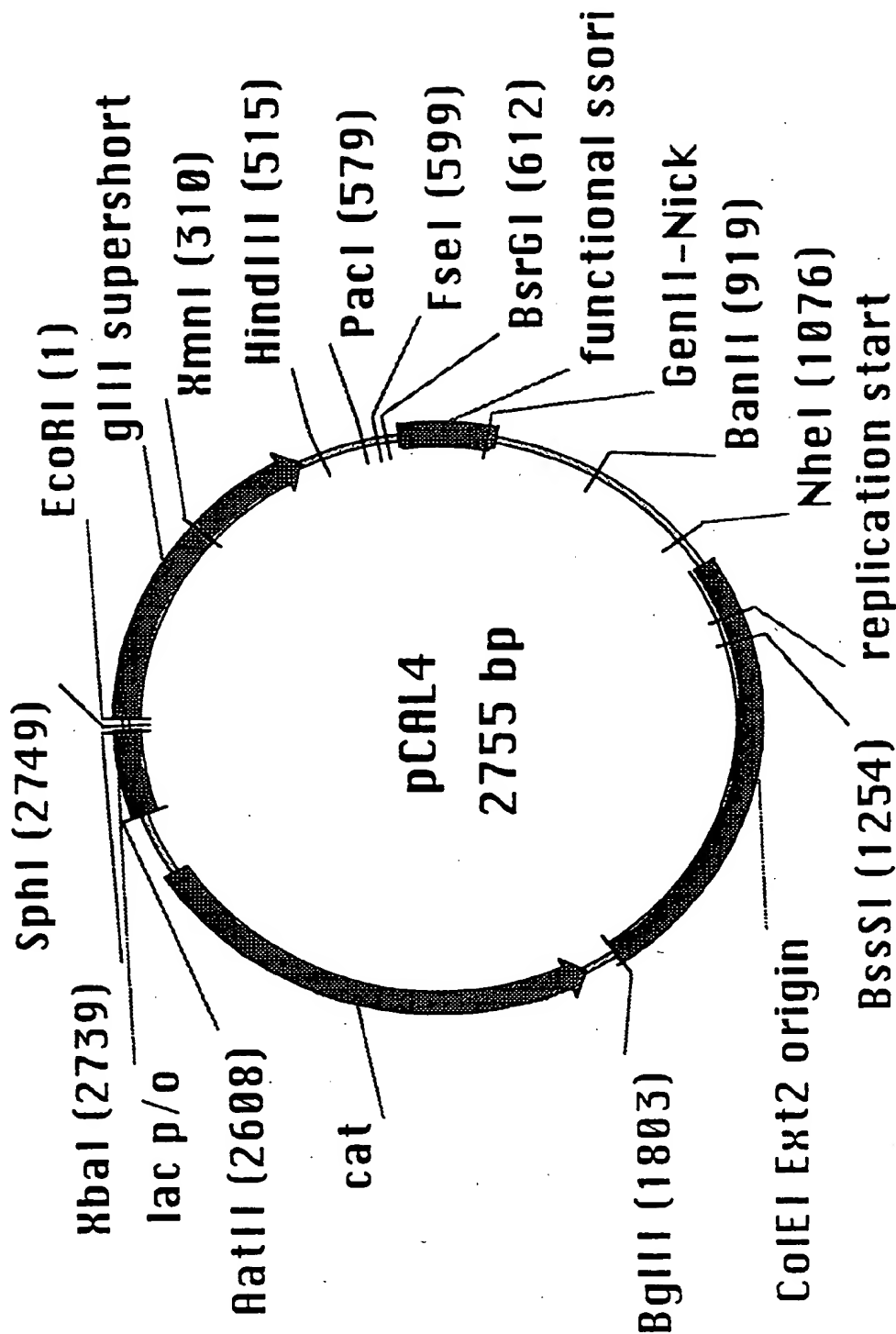


Figure 35: functional map and sequence of modular vector pCAL4

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

ECORI  
~~~~~  
1 AATTCGAGCA GAAGCTGATC TCTGAGGAGG ATCTGTAGGG TGGTGGCTCT  
TTAAGCTCGT CTTCGACTAG AGACTCCTCC TAGACATCCC ACCACCGAGA  
51 GGTTCGCGTG ATTTTGATTA TGAAAAGATG GCAAACGCTA ATAAGGGGGC  
CCAAGGCCAC TAAAACTAAT ACTTTTCTAC CGTTTGCGAT TATTCCTCCC  
101 TATGACCGAA AATGCCGATG AAAACGCGCT ACAGTCTGAC GCTAAAGGCA  
ATACTGGCTT TTACGGCTAC TTTTGCGCGA TGTCAGACTG CGATTTCCTG  
151 AACTTGATTC TGTCGCTACT GATTACGGTG CTGCTATCGA TGGTTTCATT  
TTGAACCTAAG ACAGCGATGA CTAATGCCAC GACGATAGCT ACCAAAGTAA  
201 GGTGACGTTT CCGGCCCTTGC TAATGGTAAT GGTGCTACTG GTGATTTTGC  
CCACTGCCAA GGCCGGAACG ATTACCATTG CCACGATGAC CACTAAAACG  
251 TGGCTCTAAT TCCCAAATGG CTCAAGTCGG TGACGGTGAT AATTCACCTT  
ACCGAGATTA AGGGTTTACC GAGTTCAGCC ACTGCCACTA TTAAGTGGAA  
XmnI  
~~~~~  
301 TAATGAATAA TTTCCGTCAA TATTACCTT CCCTCCCTCA ATCGGTGAA  
ATTACTTATT AAAGGCAGTT ATAAATGGAA GGGAGGGAGT TAGCCAACTT



Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|     |             |            |            |             |            |
|-----|-------------|------------|------------|-------------|------------|
| 351 | TGTCGCCCTT  | TTGTCTTTGG | CGCTGGTAAA | CCATATGAAT  | TTTCTATTGA |
|     | ACAGCGGGAA  | AACAGAAACC | CGGACCATTT | GGTATACTTA  | AAAGATAACT |
| 401 | TTGTGACAAA  | ATAAACTTAT | TCCGTGGTGT | CTTTGCCGTTT | CTTTTATATG |
|     | AACACTGTTT  | TATTTGAATA | AGGCACCACA | GAAACGCAAA  | GAAAATATAC |
| 451 | TTGCCACCTT  | TATGTATGTA | TTTTCTACGT | TTGCTAACAT  | ACTGCGTAAT |
|     | AACGGTGGAA  | ATACATACAT | AAAAGATGCA | AACGATTGTA  | TGACGCATTA |
|     |             | HindIII    |            |             |            |
|     |             | ~~~~~      |            |             |            |
| 501 | AAGGAGTCTT  | GATAAGCTTG | ACCTGTGAAG | TGAAAAATGG  | CGCAGATTGT |
|     | TTCCCTCAGAA | CTATTGGAAC | TGGACACTTC | ACTTTTTACC  | CGGTCTAACA |
|     |             |            | PacI       |             | FseI       |
|     |             |            | ~~~~~      |             | ~~~~~      |
| 551 | GGGACATTTT  | TTTTGTCTGC | CGTTTAATTA | AAGGGGGGGG  | GGGGCCGGCC |
|     | CGCTGTAAAA  | AAACAGACG  | GCAAATTAAT | TTCCCCCCCC  | CCCCGGCCGG |
|     |             |            | BsrGI      |             |            |
|     |             |            | ~~~~~      |             |            |
| 601 | TGGGGGGGGG  | TGTACATGAA | ATTGTAAACG | TTAATATTTT  | GTAAAAATTC |
|     | ACCCCCCCCC  | ACATGTACTT | TAACATTTGC | AATTATAAAA  | CAATTTTAAG |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|       |             |            |            |            |            |
|-------|-------------|------------|------------|------------|------------|
| 651   | GCGTTAAATT  | TTTGTTAAAT | CAGCTCATT  | TTTAACCAAT | AGGCCGAAAT |
|       | CGCAATTTAA  | AAACAATTAA | GTCGAGTAAA | AAATTGGTTA | TCCGGCTTTA |
| 701   | CGGCAAAATC  | CCTTATAAAT | CAAAAGAATA | GACCGAGATA | GGTTGAGTG  |
|       | GCCGTTTTAG  | GGAATATTAA | GTTTCTTAT  | CTGGCTCTAT | CCCAACTCAC |
| 751   | TTGTTCCAGT  | TTGGAACAAG | AGTCCACTAT | TAAAGAACGT | GGACTCCAAC |
|       | AACAAGGTCA  | AACCTTGTC  | TCAGGTGATA | ATTCTTGCA  | CCTGAGGTG  |
| 801   | GTCAAAGGGC  | GAAAACCCGT | CTATCAGGGC | GATGGCCCAC | TACGAGAACC |
|       | CAGTTTCCCG  | CTTTTGGCA  | GATAGTCCCG | CTACCGGGTG | ATGCTCTTGG |
| 851   | ATCACCCCTAA | TCAAGTTT   | TGGGTCGAG  | GTGCCGTAAA | GCACTAAATC |
|       | TAGTGGGATT  | AGTTCAAAAA | ACCCAGCTC  | CACGGCATTT | CGTGATTTAG |
| BanII |             |            |            |            |            |
| ~~~~~ |             |            |            |            |            |
| 901   | GGAACCCCTAA | AGGGAGCCCC | CGATTAGAG  | CTTGACGGGG | AAAGCCGGCG |
|       | CCTTGGGATT  | TCCCTCGGGG | GCTAAATCTC | GAACTGCCCC | TTTCGGCCCG |
| 951   | AACGTGGCGA  | GAAAGGAAGG | GAAGAAAGCG | AAAGGAGCGG | GCGCTAGGGC |
|       | TTGCACCGCT  | CTTTCCTTCC | CTTCTTTTCG | TTTCCTCGCC | CGCGATCCCG |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|       |            |            |             |            |            |
|-------|------------|------------|-------------|------------|------------|
| 1001  | GCTGGCAAGT | GTAGCGGTCA | CGCTGCGCGT  | AACCACCACA | CCCGCCGCGC |
|       | CGACCGTTCA | CATCGCCAGT | GCGACGCGCA  | TTGGTGGTGT | GGCGGCGCGC |
| NheI  |            |            |             |            |            |
|       | ~~~~~      |            |             |            |            |
| 1051  | TTAATGCGCC | GCTACAGGC  | GCGTGCTAGC  | CATGTGAGCA | AAAGGCCAGC |
|       | AATTACGCGG | CGATGTCCCG | GCGACGATCG  | GTACACTCGT | TTTCCGCTCG |
| 1101  | AAAAGGCCAG | GAACCGTAA  | AAGGCCGCGT  | TGCTGGCGTT | TTTCCATAGG |
|       | TTTTCGCGTC | CTTGGCATT  | TTCCGGCGCA  | ACGACCGCAA | AAAGGTATCC |
| 1151  | CTCCGCCCCC | CTGACGAGCA | TCACAAAAT   | CGACGCTCAA | GTCAGAGGTG |
|       | GAGCGGGGG  | GACTGCTCGT | AGTGTTTTTA  | GCTGCGAGTT | CAGTCTCCAC |
| 1201  | GCGAAACCCG | ACAGGACTAT | AAAGATACCA  | GGCGTTTCCC | CCTGGAAGCT |
|       | CGCTTTGGGC | TGTCCTGATA | TTTCTATGGT  | CCGCAAAGGG | GGACCTTCGA |
| BssSI |            |            |             |            |            |
|       | ~~~~~      |            |             |            |            |
| 1251  | CCCTCGTGCG | CTCTCCTGTT | CCGACCCCTGC | CGCTTACCGG | ATACCTGTCC |
|       | GGGAGCACGC | GAGAGGACAA | GGCTGGGACG  | GCGAATGGCC | TATGGACAGG |
| 1301  | GCCTTTCTCC | CTTCGGGAAG | CGTGGCGCTT  | TCTCATAGCT | CACGCTGTAG |
|       | CGGAAAGAGG | GAAGCCCTTC | GCACCGCGAA  | AGAGTATCGA | GTGCGACATC |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|      |             |             |            |             |             |
|------|-------------|-------------|------------|-------------|-------------|
| 1351 | GTATCTCAGT  | TCGGTGTAGG  | TCGTTGCTC  | CAAGCTGGGC  | TGTGTCACG   |
|      | CATAGAGTCA  | AGCCACATCC  | AGCAAGCGAG | GTTCCGACCCG | ACACACGTGC  |
| 1401 | AACCCCCCGT  | TCAGCCCCGAC | CGCTGCGCCT | TATCCGGTAA  | CTATCGTCTT  |
|      | TTGGGGGGCA  | AGTCGGGCTG  | GCGACGGGA  | ATAGGCCATT  | GATAGCAGAA  |
| 1451 | GAGTCCAACC  | CGGTAAGACA  | CGACTTATCG | CCACTGGCAG  | CAGCCACTGG  |
|      | CTCAGGTTGG  | GCCATTCTGT  | GCTGAATAGC | GGTGACCGTC  | GTCGGTGACC  |
| 1501 | TAACAGGATT  | AGCAGAGCGA  | GGTATGTAGG | CGGTGCTACA  | GAGTCTTGA   |
|      | ATTGTCCTAA  | TCGTCTCGCT  | CCATACATCC | GCCACGATGT  | CTCAAGAACT  |
| 1551 | AGTGGTGGCC  | TAACTACGGC  | TACACTAGAA | GAACAGTATT  | TGGTATCTGC  |
|      | TCACCAACCG  | ATTGATGCCG  | ATGTGATCTT | CTTGTCATAA  | ACCATAGACG  |
| 1601 | GCTCTGCTGT  | AGCCAGTTAC  | CTTCGGAAAA | AGAGTTGGTA  | GCTCTTGATC  |
|      | CGAGACGACA  | TCGGTCAATG  | GAAGCCTTTT | TCTCAACCAT  | CGAGAACTAG  |
| 1651 | CGGCAAACAA  | ACCACCGCTG  | GTAGCGGTGG | TTTTTTTGT   | TGCAAGCAGC  |
|      | GCCGTTTGT   | TGGTGGCGAC  | CATCGCCACC | AAAAAAACAA  | ACGTTTCGTG  |
| 1701 | AGATTACGGC  | CAGAAAAAAA  | GGATCTCAAG | AAGATCCTTT  | GATCTTTTCT  |
|      | TCCTAATGCGC | GTCTTTTTTT  | CCTAGAGTTC | TTCTAGGAAA  | CTAGAAAAAGA |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|      |                        |             |            |            |            |
|------|------------------------|-------------|------------|------------|------------|
| 1751 | ACGGGGTCTG             | ACGCTCAGTG  | GAACGAAAC  | TCACGTTAAG | GGATTTTGGT |
|      | TGCCCCCAGAC            | TGCGAGTCAC  | CTTGCTTTTG | AGTGCAATC  | CCTAAACCA  |
|      | <p>BglII<br/>~~~~~</p> |             |            |            |            |
| 1801 | CAGATCTAGC             | ACCAGGCGTT  | TAAGGCACC  | AATAACTGCC | TTAAAAAAT  |
|      | GTCTAGATCG             | TGGTCCGCAA  | ATTCCCGTGG | TTATTGACGG | AATTTTTTA  |
| 1851 | TACGCCCCCGC            | CCTGCCACTC  | ATCGCAGTAC | TGTTGTAATT | CATTAAGCAT |
|      | ATGCGGGGCG             | GGACGGTGAG  | TAGCGTCATG | ACAACATTAA | GTAATTGTA  |
| 1901 | TCTGCCGACA             | TGGAAGCCAT  | CACAAACGGC | ATGATGAACC | TGAATCGCCA |
|      | AGACGGCTGT             | ACCTTCGGTA  | GTGTTTGCCG | TACTACTTGG | ACTTAGCGGT |
| 1951 | GCGGCATCAG             | CACCTTGTCG  | CCTTGCGTAT | AATATTGCC  | CATAGTAAA  |
|      | CGCCGTAGTC             | GTGGAACAGC  | GGAACGCATA | TTATAAACGG | GTATCATT   |
| 2001 | ACGGGGGCGA             | AGAAGTTGTC  | CATATTGGCT | ACGTTTAAAT | CAAAACTGGT |
|      | TGCCCCCGCT             | TCTTCAACAG  | GTATAACCGA | TGCAAAATTA | GTTTGGACCA |
| 2051 | GAAACTCACC             | CAGGGATTGG  | CTGAGACGAA | AAACATATTC | TCAATAAACC |
|      | CTTTGAGTGG             | GTCCCCTAACC | GACTCTGCTT | TTTGTATAAG | AGTTATTGG  |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

|      |                                                         |                                                         |
|------|---------------------------------------------------------|---------------------------------------------------------|
| 2101 | CTTTAGGGAA ATAGGCCAGG TTTTCACCGT AACACGCCAC ATCTTGCGAA  | GAAATCCCTT TATCCGGTCC AAAAGTGGCA TTGTGCGGTG TAGAACGCTT  |
| 2151 | TATATGTGTA GAAACTGCCG GAAATCGTCG TGGTATTTCAC TCCAGAGCGA | ATATACACAT CTTTGACGGC CTTTAGCAGC ACCATAAGTG AGGTCTCGCT  |
| 2201 | TGAAAACGTT TCAGTTTGCT CATGAAAC GGTGTAACAA GGTGAACAC     | ACTTTTGCAA AGTCAAACGA GTACCTTTTG CCACATTGTT CCCACTTG TG |
| 2251 | TATCCCATAT CACCAGCTCA CCGTCTTTCA TTGCCATACG GAACTCCGGG  | ATAGGGTATA GTGGTCGAGT GGCAGAAAGT AACGGTATGC CTTGAGGCC   |
| 2301 | TGAGCATTCA TCAGCGGGC AAGAATGTGA ATAAAGGCCG GATAAACTT    | ACTCGTAAGT AGTCCGCCCG TTCTTACACT TATTTCGGC CTATTTTGAA   |
| 2351 | GTGCTTATT TTCTTTACGG TCTTTAAAAA GGCCGTAATA TCCAGCTGAA   | CACGAAATAA AAGAAATGCC AGAAATTTT CCGGCATTAT AGGTCGACTT   |
| 2401 | CGGTCTGGTT ATAGGTACAT TGAGCAACTG ACTGAAATGC CTCAAAATGT  | GCCAGACCAA TATCCATGTA ACTCGTTGAC TGACTTTACG GAGTTTAC    |
| 2451 | TCCTTACGAT GCCATTGGGA TATATCAACG GTGGTATATC CAGTGATTTT  | AGAAATGCTA CGGTAACCTT ATATAGTTGC CACCATATAG GTCACATAAA  |

Figure 35: functional map and sequence of modular vector pCAL4 (continued)

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2501 TTCTCTCCATT TTAGCTTCCT TAGCTCCTGA AAATCTCGAT AACTCAAAA
 AAAGAGGTAA AATCGAAGGA ATCGAGGACT TTTAGAGCTA TTGAGTTTTT

2551 ATACGCCCGG TAGTGATCTT ATTTCAATAT GTGAAAGTT GGAACCTCAC
 TATGCGGGCC ATCACTAGAA TAAAGTAATA CCACTTTCAA CCTTGGAGTG

 AatII
      ~~~~~

2601  CCGACGTCTA ATGTGAGTTA GCTCACTCAT TAGGCACCCC AGGCTTTACA
      GGCTGCAGAT TACACTCAAT CGAGTGAGTA ATCCGTGGGG TCCGAAATGT

2651  CTTTATGCTT CCGGCTCGTA TGTGTGTGG AATTGTGAGC GGATAACAAT
      GAAATACGAA GGCCGAGCAT ACAACACACC TTAACACTCG CCTATTGTTA

      XbaI   SphI
      ~~~~~

2701 TTCACACAGG AACAGCTAT GACCATGATT ACCAATTCT AGAGCATGCG
 AAGTGTGTCC TTTGTGCGATA CTGGTACTAA TGCTTAAAGA TCTCGTACGC

 EcoRI

2751 GGGGG
 CCCCC

```





Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 2:

|     |             |            |            |                        |
|-----|-------------|------------|------------|------------------------|
|     | AatII       |            |            |                        |
|     | ~~~~~       |            |            |                        |
| 1   | GACGTCCTTAA | TGTGAGTTAG | CTCACTCATT | AGGCACCCCA GGCTTTACAC  |
|     | CTGCAGAATT  | ACACTCAATC | GAGTGAGTAA | TCCGTGGGGT CCGAAATGTG  |
| 51  | TTATGCTTC   | CGGCTCGTAT | GTTGTGTGGA | ATTGTGAGCG GATAACAATT  |
|     | AAATACGAAG  | GCCGAGCATA | CAACACACCT | TAACTACTCGC CTATTGTTAA |
|     |             |            | XmnI       | ~~~~~                  |
|     |             |            | XbaI       | ~~~~~                  |
| 101 | TCACACAGGA  | AACAGCTATG | ACCATGTCTA | GAATAACTTC GTATAATGTA  |
|     | AGTGTGTCCT  | TTGTCGATAC | TGGTACAGAT | CTTATTGAAG CATATTACAT  |
|     |             |            | SphI       | ~~~~~                  |
| 151 | CGCTATACGA  | AGTATCGCA  | TGC        |                        |
|     | GCGATATGCT  | TCAATAGCGT | ACG        |                        |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

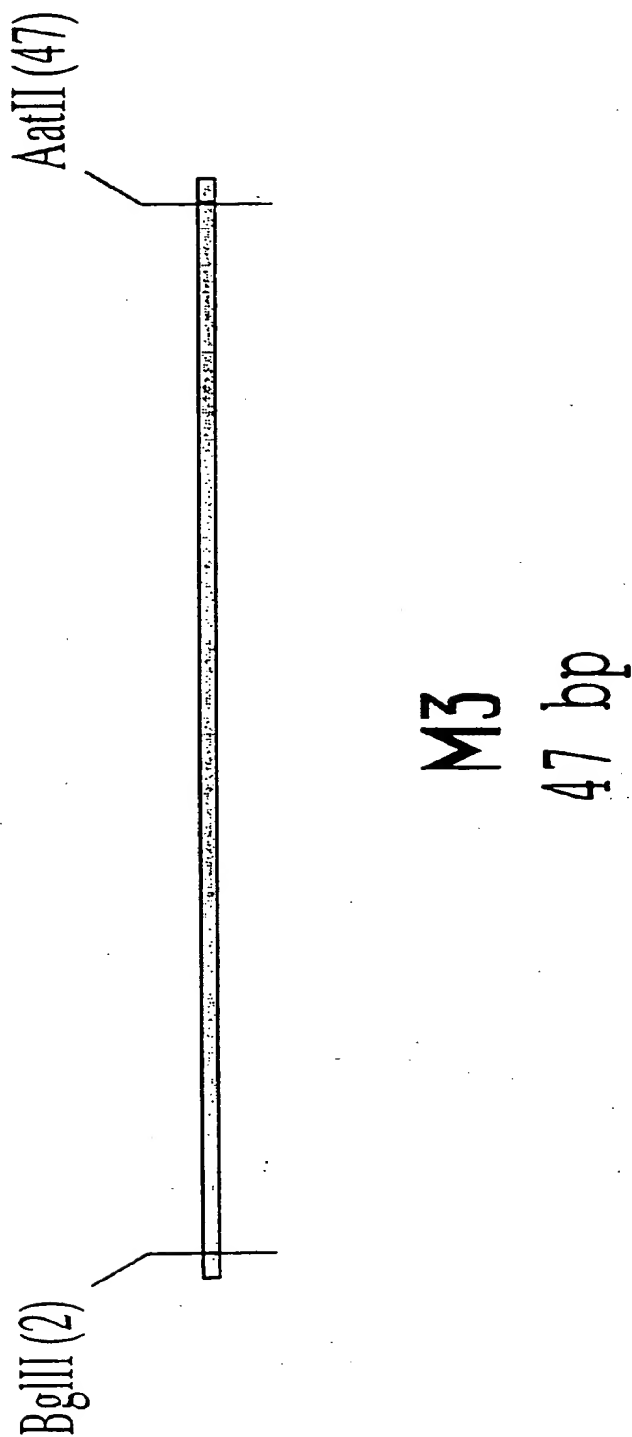


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 3:

|   |                                                     |  |       |
|---|-----------------------------------------------------|--|-------|
|   | BglII                                               |  | AatII |
|   | ~~~~~                                               |  | ~~~~~ |
| 1 | AGATCTCATA ACTTCGTATA ATGTATGCTA TACGAAGTTA TGACGTC |  |       |
|   | TCTAGAGTAT TGAAGCATAT TACATACGAT ATGCTTCAAT ACTGCAG |  |       |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

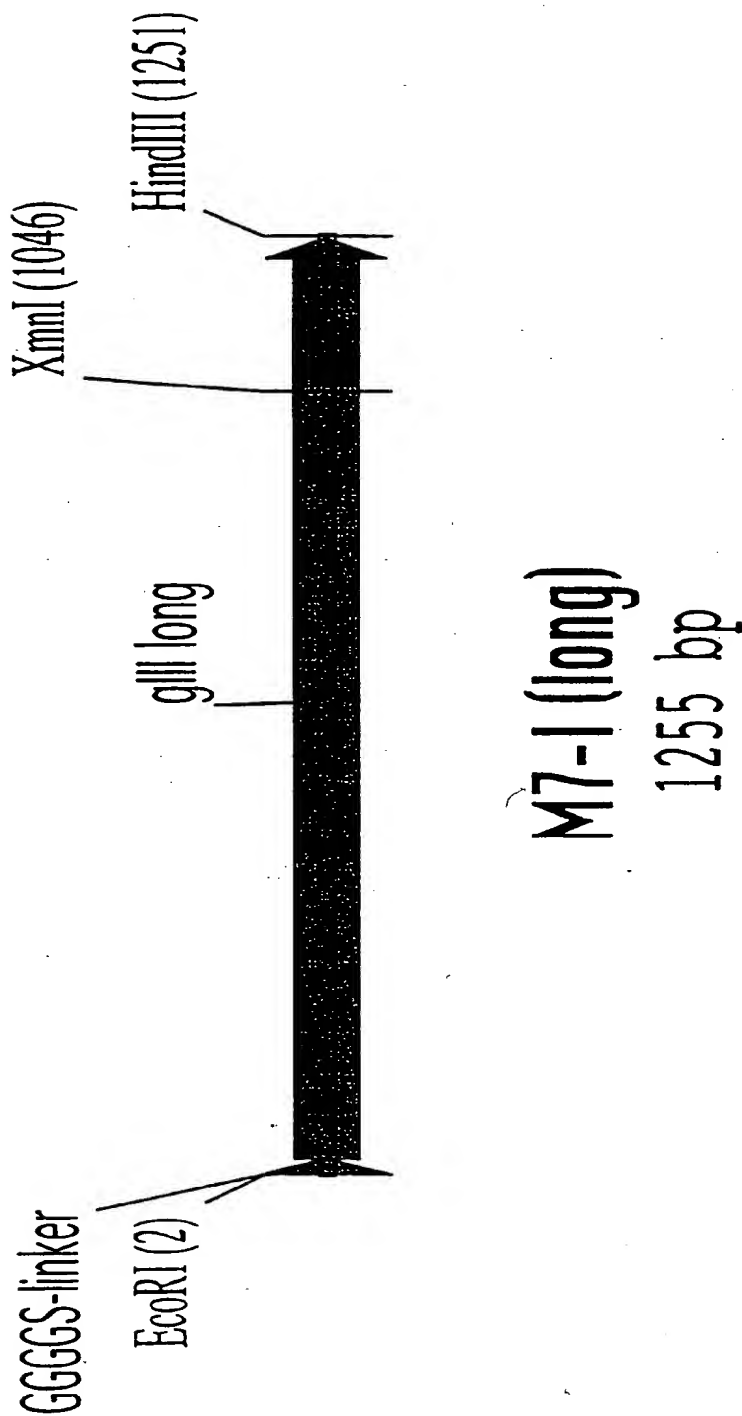


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 7-I (long):

ECORI

-----

|     |            |             |             |            |             |
|-----|------------|-------------|-------------|------------|-------------|
| 1   | GAATTCGGTG | GTGGTGGATC  | TGCGTGCGCT  | GAAACGGTTG | AAAGTTGTTT  |
|     | CTTAAGCCAC | CACCACCTAG  | ACGCACGCGA  | CTTTGCCAAC | TTTCAACAAA  |
| 51  | AGCAAAATCC | CATACAGAAA  | ATTCAATTAC  | TAACGTCTGG | AAAGACGACA  |
|     | TCGTTTTAGG | GTATGTCTTT  | TAAAGTAAATG | ATTGCAGACC | TTTCTGCTGT  |
| 101 | AAACTTTAGA | TCGTTACGCT  | AACATAGAGG  | GCTGCTCTGT | GAATGCTACA  |
|     | TTTGAAATCT | AGCAATGCGA  | TTGATACTCC  | CGACAGACAC | CTTACGATGT  |
| 151 | GGCGTTGTAG | TTTGTA CTGG | TGACGAAACT  | CAGTGTTACG | GTACATGGGT  |
|     | CCGCAACATC | AAACATGACC  | ACTGCTTTGA  | GTCACAATGC | CATGTACCCA  |
| 201 | TCCTATTGGG | CTTGCTATCC  | CTGAAAATGA  | GGGTGGTGGC | TCTGAGGGTG  |
|     | AGGATAACCC | GAACGATAGG  | GACTTTTACT  | CCCACCCACG | AGACTCCCCAC |
| 251 | GCGGTTCTGA | GGGTGGCGGT  | TCTGAGGGTG  | GCGGTACTAA | ACCTCCTGAG  |
|     | CGCCAAGACT | CCCACCGCCA  | AGACTCCCCAC | CGCCATGATT | TGGAGGACTC  |
| 301 | TACGGTGATA | CACCTATTCC  | GGGCTATACT  | TATATCAACC | CTCTCGACGG  |
|     | ATGCCACTAT | GTGGATAAGG  | CCCGATATGA  | ATATAGTTGG | GAGAGCTGCC  |

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |             |             |            |             |             |
|-----|-------------|-------------|------------|-------------|-------------|
| 351 | CACTTATCCG  | CCTGGTACTG  | AGCAAAACCC | CGCTAATCCT  | AATCCTTCTC  |
|     | GTGAATAGGC  | GGACCATGAC  | TCGTTTGGG  | GCGATTAGGA  | TTAGGAAGAG  |
| 401 | TTGAGGAGTC  | TCAGCCCTCTT | AATACTTCA  | TGTTTCAGAA  | TAATAGGTTC  |
|     | AACTCCTCAG  | AGTCGGAGAA  | TTATGAAAGT | ACAAAGTCTT  | ATTATCCAAG  |
| 451 | CGAAATAGGC  | AGGGGGCATT  | AACTGTTTAT | ACGGGCACTG  | TTACTCAAGG  |
|     | GCTTTATCCG  | TCCCCCGTAA  | TTGACAAATA | TGCCCCGTGAC | AATGAGTTCC  |
| 501 | CACTGACCCC  | GTTAAACTT   | ATTACCAGTA | CACTCCTGTA  | TCATCAAAAG  |
|     | GTGACTGGGG  | CAATTTTGAA  | TAATGGTCAT | GTGAGGACAT  | AGTAGTTTTC  |
| 551 | CCATGTATGA  | CGCTTACTGG  | AACGGTAAAT | TCAGAGACTG  | CGCTTTCCAT  |
|     | GGTACATACT  | GCGAATGACC  | TTGCCATTTA | AGTCTCTGAC  | GCGAAAGGTA  |
| 601 | TCTGGCTTTA  | ATGAGGATTT  | ATTGTGTTGT | GAATATCAAG  | GCCAATCGTC  |
|     | AGACCGAAAT  | TACTCCCTAAA | TAAACAAACA | CTTATAGTTC  | CGGTTAGCAG  |
| 651 | TGACCTGCCT  | CAACCTCCTG  | TCAATGCTGG | CGGCGGCTCT  | GGTGGTGGTT  |
|     | ACTGGACGGA  | GTTGGAGGAC  | AGTTACGACC | GCCGCCGAGA  | CCACCACCAA  |
| 701 | CTGGTGCGCG  | CTCTGAGGGT  | GGTGGCTCTG | AGGGTGGCGG  | TTCTGAGGGT  |
|     | GACCAACCGCC | GAGACTCCCCA | CCACCGAGAC | TCCCACCGCC  | AAGACTCCCCA |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |            |            |            |            |             |
|------|------------|------------|------------|------------|-------------|
| 751  | GGCGGCTCTG | AGGAGGCGG  | TTCGGGTGGT | GGCTCTGGTT | CCGGTGATTT  |
|      | CCGCCGAGAC | TCCCTCCGCC | AAGGCCACCA | CCGAGACCAA | GGCCACTAAA  |
| 801  | TGATTATGAA | AAGATGGCAA | ACGCTAATAA | GGGGGCTATG | ACCGAAAATG  |
|      | ACTAATACTT | TTCTACCGTT | TCCGATTATT | CCCCCGATAC | TGGCTTTTAC  |
| 851  | CCGATGAAAA | CGCGCTACAG | TCTGACGCTA | AAGGCAAACT | TGATTCTGTC  |
|      | GGCTACTTTT | CGCGGATGTC | AGACTGCGAT | TTCCGTTTGA | ACTAAGACAG  |
| 901  | GCTACTGATT | ACGGTGCTGC | TATCGATGGT | TTCATTGGTG | ACGTTTCCGG  |
|      | CGATGACTAA | TGCCACGACG | ATAGCTACCA | AAGTAACCAC | TGCAAAAGGCC |
| 951  | CCTTGCTAAT | GGTAATGGTG | CTACTGGTGA | TTTTGCTGGC | TCTAATTCCC  |
|      | GGAACGATTA | CCATTACCAC | GATGACCACT | AAAACGACCG | AGATTAAAGG  |
| 1001 | AAATGGCTCA | AGTCGGTGAA | GGTGATAAAT | CACCTTTAAT | GAATAATTTC  |
|      | TTTACCGAGT | TCAGCCACTT | CCACTATTAA | GTGGAAATTA | CTTATTAAAG  |
| 1051 | CGTCAATATT | TACCTTCCAT | CCCTCAATCG | GTGAAATGTC | GCCCTTTTGT  |
|      | GCAGTTATAA | ATGGAAGGTA | GGGAGTTAGC | CAACTTACAG | CGGAAAACAA  |

XmnI

~~~~~

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |            |             |            |            |            |
|------|------------|-------------|------------|------------|------------|
| 1101 | CTTTGGCGCT | GGTAAACCCCT | ATGAATTTTC | TATTGATTGT | GACAAAATAA |
|      | GAAACCGCGA | CCATTGGGA   | TACTTAAAG  | ATAACTAACA | CTGTTTTATT |
| 1151 | ACTTATTCCG | TGGTGCTTT   | GCGTTTCTTT | TATATGTTGC | CACCTTTATG |
|      | TGAATAAGGC | ACCACAGAAA  | CGCAAAGAAA | ATATACAACG | GTGGAAATAC |
|      |            |             |            |            | HindIII    |
| 1201 | TATGTATTTT | CTACGTTTGC  | TAACATACTG | CGTAATAAGG | AGTCTTGATA |
|      | ATACATAAAA | GATGCAAACG  | ATTGTATGAC | GCATTATTCC | TCAGAACTAT |

HindI  
 ~~~  
 AGCTT  
 TCGAA

1251



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

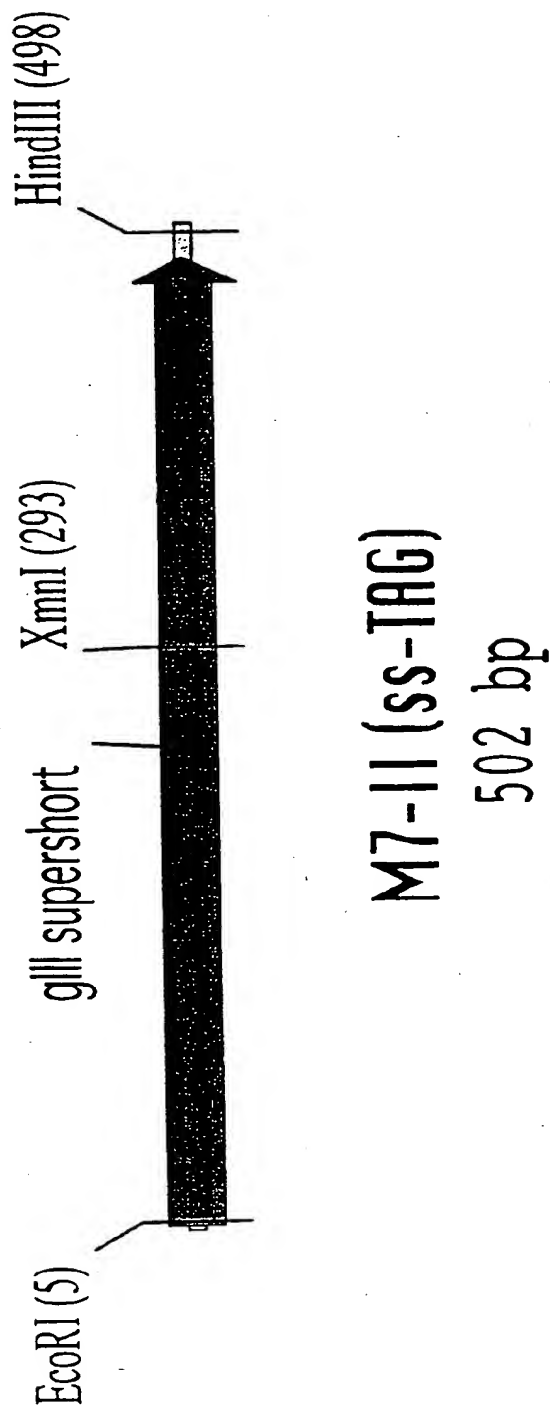
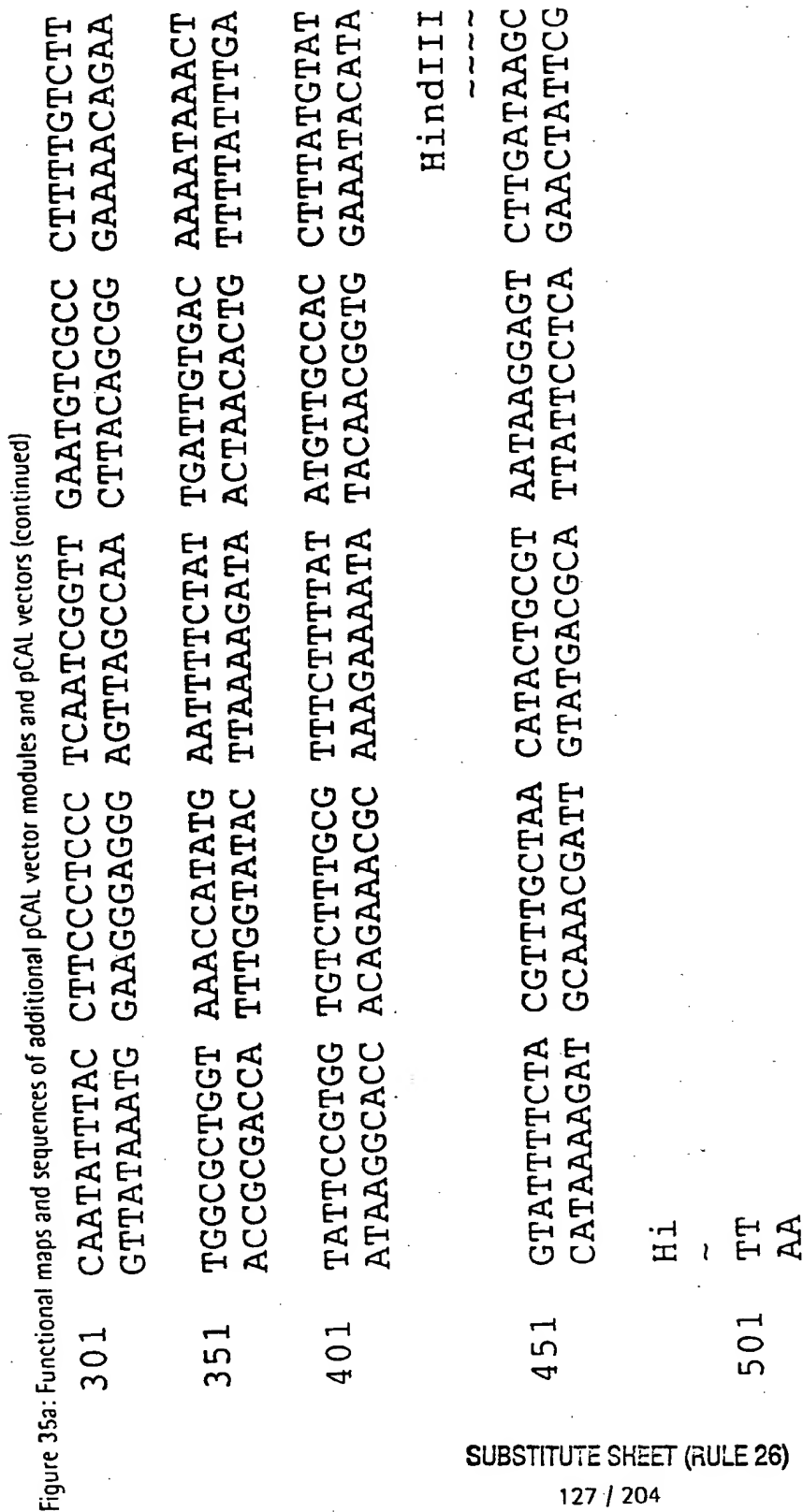


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 7-II (SS-TAG):

| ECORI |                                                                                                                    |
|-------|--------------------------------------------------------------------------------------------------------------------|
| ~~~~~ |                                                                                                                    |
| 1     | CGGGAATTTC GAGCGGGTTC CGGTGGTGGC TCTGGTTCCG GTGATTTTGA<br>GCCCTTAAGC CTCGGCCAAG GCCACCACCG AGACCAAGGC CACTAAAACT   |
| 51    | TTATGAAAAG ATGGCAAACG CTAATAAGGG GGCTATGACC GAAAATGCCG<br>AATACTTTTC TACCGTTTGC GATTATTCCC CCGATACTGG CTTTTACGGC   |
| 101   | ATGAAAACGC GCTACAGTCT GACGCTAAAG GCAAAC TTGA TTCTGTCGCT<br>TACTTTTGCG CGATGTCAGA CTGCGATTTC CGTTTGA ACT AAGACAGCGA |
| 151   | ACTGATTACG GTGCTGCTAT CGATGGTTTC ATTGGTGACG TTTCCGGCCT<br>TGACTAATGC CACGACGATA GCTACCAAAG TAACCACTGC AAAGGCCGGA   |
| 201   | TGCTAATGGT AATGGTGCTA CTGGTGATT TGCTGGCTCT AATCCCAA<br>ACGATTACCA TTACCACGAT GACCACTAAA ACGACCGAGA TTAAGGGTTT      |
| XmnI  |                                                                                                                    |
| ~~~~~ |                                                                                                                    |
| 251   | TGGCTCAAGT CCGTGACGGT GATAATTCAC CTTTAATGAA TAATTTCCGT<br>ACCGAGTTCA GCCACTGCCA CTATTAAGTG GAAATTACTT ATTAAGGCA    |



SphI (6)

$$\frac{\infty}{\Sigma}$$

47 bp

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 8:

|   |                                                      |  |         |
|---|------------------------------------------------------|--|---------|
|   | SphI                                                 |  | HindIII |
|   | -----                                                |  | -----   |
| 1 | GCATGCCATA ACTTCGTATA ATGTACGCTA TACGAAGTTA TAAGCTT  |  | TAAGCTT |
|   | CGTACGGTAT TGAAGCATAT TACATGCCGAT ATGCTTCAAT ATTCGAA |  | ATTCGAA |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

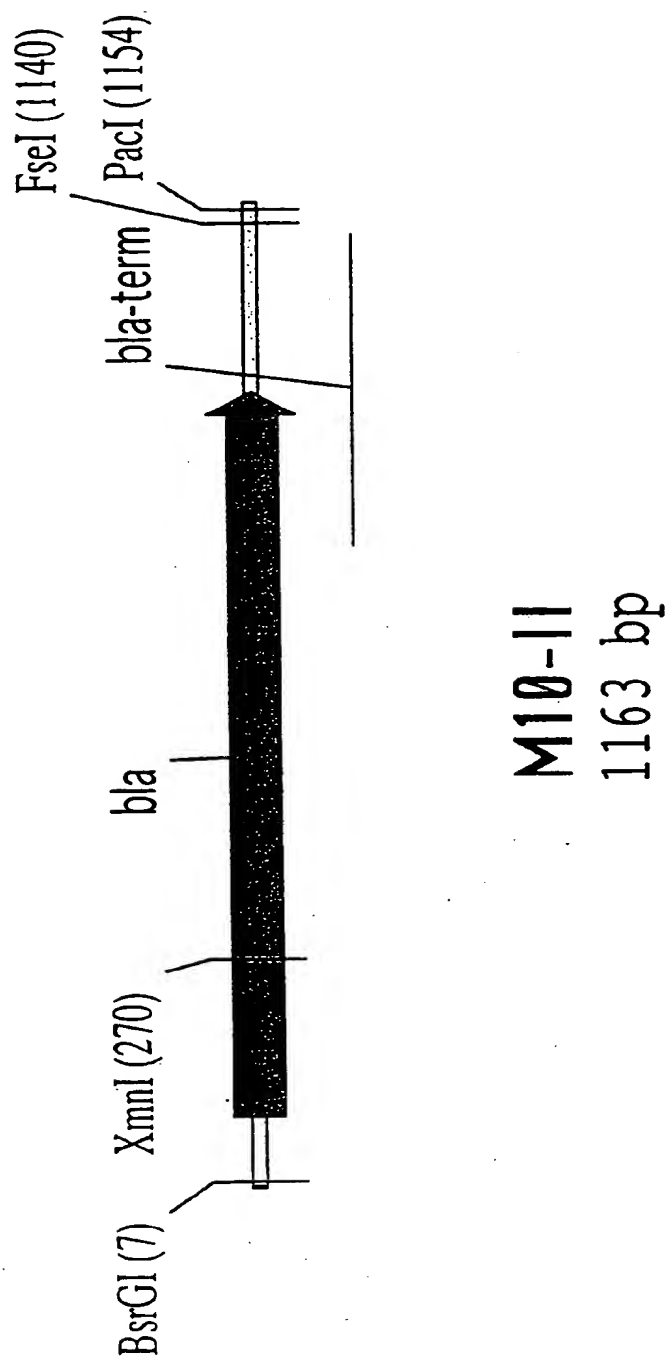


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 10-II:

BsrGI

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1 GGGGGTGTAC ATTCAAATAT GTATCCGCTC ATGAGACAAT AACCTGATA
 CCCCACATG TAAGTTTATA CATAGGCGAG TACTCTGTTA TTGGGACTAT

51 AATGCTTCAA TAATATTGAA AAAGGAAGAG TATGAGTATT CAACATTTC
 TTACGAAGTT ATTATAACTT TTTCCCTTCTC ATACTCATAA GTTGTAAGG

101 GTGTCGCCCT TATTCCTTT TTTGCGGCAT TTTGCCCTCC TGTTTTGCT
 CACAGCGGGA ATAAGGGAAA AAACGCCGTA AAACGGAAGG ACAAAAACGA

151 CACCCAGAAA CGCTGGTGAA AGTAAAAGAT GCTGAGGATC AGTTGGGTGC
 GTGGGTCTTT GCGACCACTT TCATTTTCTA CGACTCCCTAG TCAACCCACG

201 GCGAGTGGGT TACATCGAAC TGGATCTCAA CAGCGGTAAG ATCCTTGAGA
 CGCTCACCCA ATGTAGCTTG ACCTAGAGTT GTCGCCATTG TAGGAACCTCT

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XmnI

~~~~~

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251 GTTTTCGCCC CGAAGAACGT TTTCCAATGA TGAGCACTTT TAAAGTTCTG
 CAAAAGCGGG GCTTCTTGCA AAAGGTTACT ACTCGTGAAA ATTTCAAGAC

```

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |            |             |             |             |             |
|-----|------------|-------------|-------------|-------------|-------------|
| 301 | CTATGTGGCG | CGGTATTATC  | CCGTATTGAC  | GCCGGGCAAG  | AGCAACTCGG  |
|     | GATACACCCG | GCCATAATAG  | GGCATAACTG  | CGGCCCGTTC  | TCGTTGAGCC  |
| 351 | TCGCCGCATA | CACTATTCTC  | AGAAAGACTT  | GGTTGAGTAC  | TCACCAGTCA  |
|     | AGCGCGGTAT | GTGATAAGAG  | TCTTACTGAA  | CCAACCTCATG | AGTGGTCAGT  |
| 401 | CAGAAAGCA  | TCTTACGGAT  | GGCATGACAG  | TAAGAGAATT  | ATGCAGTGCT  |
|     | GTCCTTTCCG | AGAAATGCCTA | CCGTACTGTC  | ATTCTCTTAA  | TACGTCACGA  |
| 451 | GCCATAACCA | TGAGTGATAA  | CACCTGGGCC  | AACTTACTTC  | TGACAAACGAT |
|     | CGGTATTGGT | ACTCACTATT  | GTGACGCCCG  | TTGAATGAAG  | ACTGTTGCTA  |
| 501 | CGGAGGACCG | AAGGAGCTAA  | CCGCTTTTTT  | GCACAACATG  | GGGGATCATG  |
|     | GCCTCCTGGC | TTCCCTCGATT | GGCGAAAAAA  | CGTGTTGTAC  | CCCCTAGTAC  |
| 551 | TAACTCGCCT | TGATCGTTGG  | GAACCGGAGC  | TGAATGAAGC  | CATACCAAAC  |
|     | ATTGAGCCGA | ACTAGCAACC  | CTTGGCCCTCG | ACTTACTTCG  | GTATGGTTTG  |
| 601 | GACGAGCGTG | ACACCACGAT  | GCCTGTAGCA  | ATGGCAACAA  | CGTTGCCGAA  |
|     | CTGCTCGCAC | TGTGGTGCTA  | CGGACATCGT  | TACCGTTGTT  | GCAACGCGTT  |
| 651 | ACTATTAACT | GGCGAACTAC  | TTACTCTAGC  | TTCCCGGCAA  | CAGTTAATAG  |
|     | TGATAATTGA | CCGCTTGATG  | AATGAGATCG  | AAGGGCCGTT  | GTCAATTATC  |

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |                                                         |      |                                                         |
|------|---------------------------------------------------------|------|---------------------------------------------------------|
| 701  | ACTGGATGGA GCGGATAAA GTTGCAGGAC CACTTCTGCG CTCGGCCCTT   | 701  | ACTGGATGGA GCGGATAAA GTTGCAGGAC CACTTCTGCG CTCGGCCCTT   |
|      | TGACCTACCT CCGCCTATT CAACGTCCCTG GTGAAGACGC GAGCCGGGAA  |      | TGACCTACCT CCGCCTATT CAACGTCCCTG GTGAAGACGC GAGCCGGGAA  |
| 751  | CCGGCTGGCT GGTTTATTGC TGATAAATCT GGAGCCGGTG AGCGTGGGTC  | 751  | CCGGCTGGCT GGTTTATTGC TGATAAATCT GGAGCCGGTG AGCGTGGGTC  |
|      | GGCCGACCGA CCAAATAACG ACTATTTAGA CCTCGGCCAC TCGACCCACG  |      | GGCCGACCGA CCAAATAACG ACTATTTAGA CCTCGGCCAC TCGACCCACG  |
| 801  | TCGCGGTATC ATTGCAGCAC TGGGGCCAGA TGGTAAGCCC TCCCGTATCG  | 801  | TCGCGGTATC ATTGCAGCAC TGGGGCCAGA TGGTAAGCCC TCCCGTATCG  |
|      | AGCGCCATAG TAACGTCTGT ACCCCGGTCT ACCATTCTGGG AGGCATAGC  |      | AGCGCCATAG TAACGTCTGT ACCCCGGTCT ACCATTCTGGG AGGCATAGC  |
| 851  | TAGTTATCTA CACGACGGGG AGTCAGGCAA CTATGGATGA ACGAAATAGA  | 851  | TAGTTATCTA CACGACGGGG AGTCAGGCAA CTATGGATGA ACGAAATAGA  |
|      | ATCAATAGAT GTGCTGCCCC TCAGTCCGTT GATACCTACT TGCTTTATCT  |      | ATCAATAGAT GTGCTGCCCC TCAGTCCGTT GATACCTACT TGCTTTATCT  |
| 901  | CAGATCGCTG AGATAGGTGC CTCACTGATT AAGCATTTGGG TAACTGTCAG | 901  | CAGATCGCTG AGATAGGTGC CTCACTGATT AAGCATTTGGG TAACTGTCAG |
|      | GCTAGCCGAC TCTATCCACG GAGTGACTAA TTCGTAACCC ATTGACAGTC  |      | GCTAGCCGAC TCTATCCACG GAGTGACTAA TTCGTAACCC ATTGACAGTC  |
| 951  | ACCAAGTTA CTCATATATA CTTTAGATTG ATTTAAAACT TCATTTTTAA   | 951  | ACCAAGTTA CTCATATATA CTTTAGATTG ATTTAAAACT TCATTTTTAA   |
|      | TGGTTCAAAT GAGTATATAT GAAATCTAAC TAAATTTTGA AGTAAAAATT  |      | TGGTTCAAAT GAGTATATAT GAAATCTAAC TAAATTTTGA AGTAAAAATT  |
| 1001 | TTTAAAAGGA TCTAGGTGAA GATCCTTTT GATAATCTCA TGACCAAAT    | 1001 | TTTAAAAGGA TCTAGGTGAA GATCCTTTT GATAATCTCA TGACCAAAT    |
|      | AAATTTTCCT AGATCCACTT CTAGGAAAAA CTATTAGAGT ACTGGTTTA   |      | AAATTTTCCT AGATCCACTT CTAGGAAAAA CTATTAGAGT ACTGGTTTA   |
| 1051 | CCCTTAACGT GAGTTTTCGT TCCACTGAGC GTCAGACCCC GTAGAAAAGA  | 1051 | CCCTTAACGT GAGTTTTCGT TCCACTGAGC GTCAGACCCC GTAGAAAAGA  |
|      | GGGAATTGCA CTCAAAAGCA AGTGACTCG CAGTCTGGGG CATCTTTTCT   |      | GGGAATTGCA CTCAAAAGCA AGTGACTCG CAGTCTGGGG CATCTTTTCT   |

|      | FseI                                                   | PacI |
|------|--------------------------------------------------------|------|
| 1101 | TCAAAGGATC TTCTTGAGAT CCTTTTGAT AATGGCCGGC CCCCCCCTT   |      |
|      | AGTTTCCCTAG AAGAACTCTA GGAATAACTA TTACCGGCCG GGGGGGGAA |      |

1151  
PacI  
~~~~~  
AATTAAGGG GGG  
TTAATTCCCC CCC

BsrGI (466)



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M11-II:

|     | NheI                                                                                                             |  |
|-----|------------------------------------------------------------------------------------------------------------------|--|
|     | -----                                                                                                            |  |
| 1   | GCTAGCACGC GCCCTGTAGC GCGGCATTAA GCGCGGCGGG TGTGGTGGTT<br>CGATCGTGCG CGGGACATCG CCGCGTAATT CCGCGCGCCC ACACCACCAA |  |
| 51  | ACGCGCAGCG TGACCGCTAC ACTTGCCAGC GCCCTAGCGC CCGCTCCTTT<br>TGCGCGTCGC ACTGGCGATG TGAACGGTCG CGGGATCGCG GCGAGGAAA  |  |
| 101 | CGCTTCTTC CCTTCCTTC TCGCCACGTT CGCCGGCTTT CCCCGTCAAG<br>GCGAAAGAAG GGAAGGAAAG AGCGGTGCAA GCGGCCGAAA GGGCAGTTC    |  |
|     | BanII                                                                                                            |  |
|     | -----                                                                                                            |  |
| 151 | CTCTAAATCG GGGCTCCCT TTAGGGTCC GATTAGTGC TTTACGGCAC<br>GAGATTAGC CCGGAGGGA AATCCCAAGG CTAAATCAGG AAATGCCCGTG     |  |
| 201 | CTCGACCCCA AAAAATTGA TTAGGGTGAT GGTCTCTGTA GTGGGCCATC<br>GAGCTGGGT TTTTGAAC TATCCCACTA CCAAGAGCAT CACCCGGTAG     |  |
| 251 | GGCCTGATAG ACGGTTTTTC GCCCTTTGAC GTTGGAGTCC ACGTTCTTTA<br>CGGGACTATC TGCCAAAAG CGGGAAACTG CAACCTCAGG TGCAAGAAAT  |  |

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |             |            |            |            |            |
|-----|-------------|------------|------------|------------|------------|
| 301 | ATAGTGGACT  | CTTGTTCCAA | ACTGGAACAA | CACTCAACCC | TATCTCGGTC |
|     | TATCACCTGA  | GAACAAGGTT | TGACCTTGTT | GTGAGTTGGG | ATAGAGCCAG |
| 351 | TATTCTTTTG  | ATTATAAAGG | GATTTTGCCG | ATTTCGGCCT | ATTGGTTAAA |
|     | ATAAGAAAC   | TAAATATTCC | CTAAACGGC  | TAAAGCCGGA | TAACCAATTT |
| 401 | AAATGAGCTG  | ATTAAACAAA | AATTTAACGC | GAATTTTAAC | AAAAATATAA |
|     | TTTACTCGAC  | TAAATTGTTT | TTAAATTGCG | CTTAAAAATG | TTTTATAAAT |
| 451 | CGTTTACAAT  | TTCATGTACA |            |            |            |
|     | GCAAAATGTTA | AAGTACATGT |            |            |            |

BsrGI

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

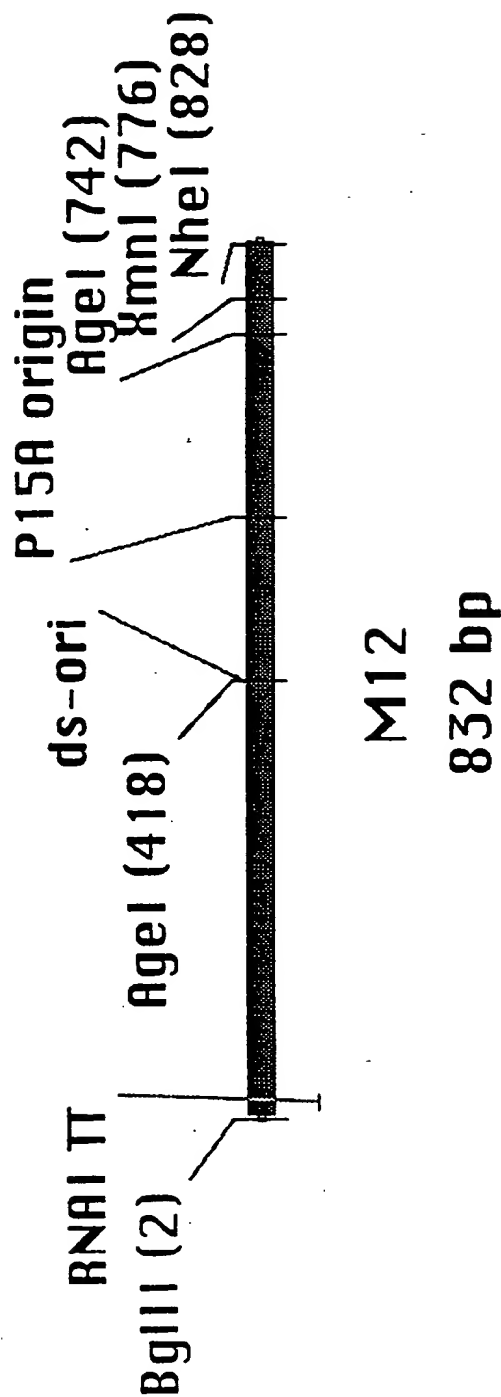


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|       |            |            |             |
|-------|------------|------------|-------------|
| M 12: |            | BglII      |             |
|       |            | ~~~~~      |             |
| 1     | AGATCTAATA | AGATGATCTT | CTTGAGATCG  |
|       | TCTAGATTAT | TCTACTAGAA | GAACCTCTAGC |
|       |            |            | AAAACCCAGAC |
|       |            |            | GCGCATTAGA  |
| 51    | CTTGCTCTGA | AAACGAAAAA | ACCGCCTTGC  |
|       | GAACGAGACT | TTTGCTTTTT | TGGCGGAACG  |
|       |            |            | TCCCGCCAAA  |
|       |            |            | AAGCATCCAA  |
|       |            |            | TTCTGTAGGTT |
| 101   | CTCTGAGCTA | CCAACCTCTT | GAACCGAGGT  |
|       | GAGACTCGAT | GGTTGAGAAA | CTTGCTCTCA  |
|       |            |            | TTGACCCGAAC |
|       |            |            | CTCCTCGCGT  |
| 151   | GTCACATAAA | CTTGTCCTTT | CAGTTAGCC   |
|       | CAGTGATTTT | GAACAGGAAA | GTCAAATCGG  |
|       |            |            | AATTGGCCGC  |
|       |            |            | CATGACTTCA  |
| 201   | AGACTAACTC | CTCTAAATCA | ATTACCAGTG  |
|       | TCTGATTGAG | GAGATTTAGT | TAATGGTCAC  |
|       |            |            | CGACGACGGT  |
|       |            |            | CACCACGAAA  |
| 251   | TGCATGTCTT | TCCGGGTTGG | ACTCAAGACG  |
|       | ACGTACAGAA | AGGCCCAACC | TGAGTTCTGC  |
|       |            |            | TATCAATGGC  |
|       |            |            | CTATTCCGCG  |
| 301   | AGCGGTCGGA | CTGAACGGGG | GGTTCGTGCA  |
|       | TCGCCAGCCT | GACTTGCCCC | CCAAGCACGT  |
|       |            |            | ATGTCAGGTC  |
|       |            |            | CTTGGAGCGA  |
|       |            |            | GAACCTCGCT  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

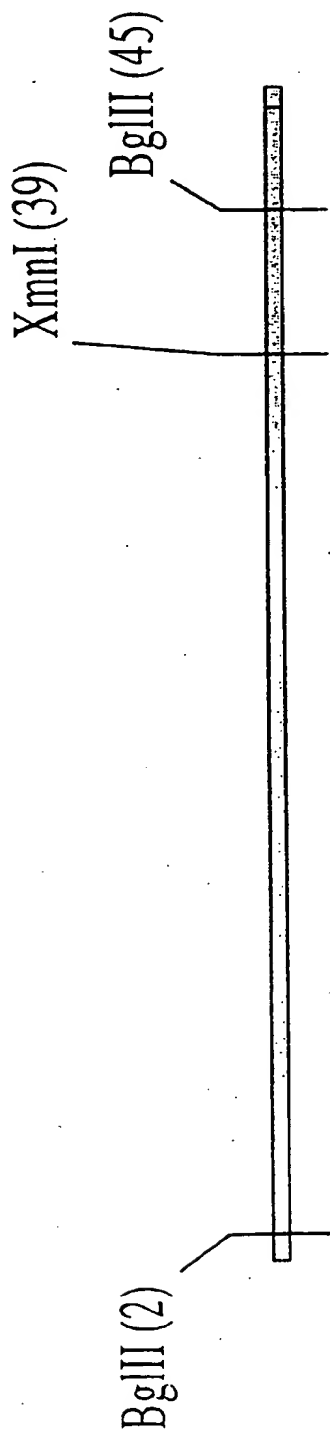
|      |             |             |             |            |             |
|------|-------------|-------------|-------------|------------|-------------|
| 351  | ACTGCCCTACC | CGGAACCTGAG | TGTCAGGCGGT | GGAATGAGAC | AAACGCGGCC  |
|      | TGACGGATGG  | GCCTTGAATC  | ACAGTCCGCA  | CCTTACTCTG | TTTGCGCGCCG |
| AgeI |             |             |             |            |             |
| 401  | ATAACAGCGG  | AATGACACCG  | GTAACCCGAA  | AGGCAGGAAC | AGGAGAGCGC  |
|      | TATTGTCGCC  | TTACTGTGGC  | CATTGGCTT   | TCCGTCCTTG | TCCTCTCGCG  |
| 451  | AGGAGGGAGC  | CGCCAGGGGG  | AAACGCCCTGG | TATCTTTATA | GTCCGTGTCGG |
|      | TCCTCCCTCG  | GCGGTCCCCC  | TTTGCGGACC  | ATAGAAATAT | CAGGACAGCC  |
| 501  | GTTTCGCCAC  | CACGTATTG   | AGCGTCAGAT  | TTCGTGATGC | TTGTCAGGGG  |
|      | CAAAGCGGTG  | GTGACTAAAC  | TCGCAGTCTA  | AAGCACTACG | AACAGTCCCC  |
| 551  | GGCGGAGCCT  | ATGGAAAAAC  | GGCTTTGCCG  | CGGCCCTCTC | ACTTCCCTGT  |
|      | CCGCCCTCGA  | TACCTTTTGG  | CCGAAACGGC  | GCCGGGAGAG | TGAAGGGACA  |
| 601  | TAAGTATCTT  | CCTGGCATCT  | TCCAGGAAAT  | CTCCGCCCCG | TTCGTAAGCC  |
|      | ATTCATAGAA  | GGACCGTAGA  | AGGTCCCTTA  | GAGCGGGGC  | AAGCATTCGG  |
| 651  | ATTTCCGCTC  | GCCGCAGTCG  | AACGACCGAG  | CGTAGCGAGT | CAGTGAGCGA  |
|      | TAAAGGCGAG  | CGCGGTCAGC  | TTGCTGGCTC  | GCATCGCTCA | GTCACCTCGCT |



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |             |            |            |            |            |
|-----|-------------|------------|------------|------------|------------|
|     |             |            |            | AgeI       |            |
|     |             |            |            | ~~~~~      |            |
| 701 | GGAAGCGGAA  | TATATCCTGT | ATCACATATT | CTGCTGACGC | ACCGGTGCAG |
|     | CCTTCGCCCTT | ATATAGGACA | TAGTGTATAA | GACGACTGCG | TGGCCACGTC |
|     |             |            |            |            |            |
|     |             |            | XmnI       |            |            |
|     |             |            | ~~~~~      |            |            |
| 751 | CCTTTTCT    | CCTGCCACAT | GAAGCACTTC | ACTGACACCC | TCATCAGTGC |
|     | GGAAAAAAGA  | GGACGGTGTA | CTTCGTGAAG | TGACTGTGGG | AGTAGTCACG |
|     |             |            |            |            |            |
|     |             |            | NheI       |            |            |
|     |             |            | ~~~~~      |            |            |
| 801 | CAACATAGTA  | AGCCAGTATA | CACTCCGCTA | GC         |            |
|     | GTTGTATCAT  | TCGGTCATAT | GTGAGGCGAT | CG         |            |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)



**M13**  
49 bp

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 13:

|   | BglII                                                 | XmnI  | BglII |
|---|-------------------------------------------------------|-------|-------|
|   | ~~~~~                                                 | ~~~~~ | ~~~~~ |
| 1 | AGATCTCATA ACTTCGTATA ATGTATGCTA TACGAAGTTA TTCAGATCT |       |       |
|   | TCTAGAGTAT TGAAGCATAT TACATACGAT ATGCTTCAAT AAGTCTAGA |       |       |

M19 96 bq 96

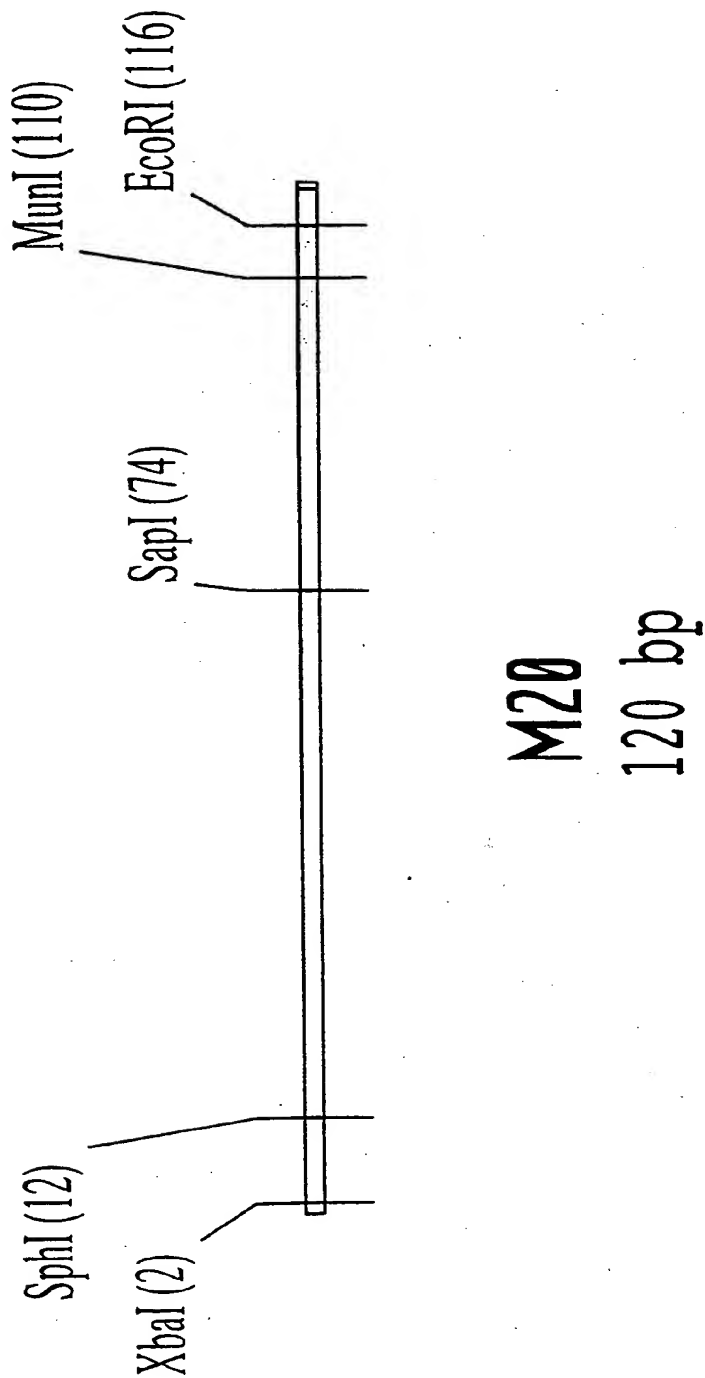
Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 19:

|    | XbaI       | SphI        |            | SapI       |            | EcoRI |
|----|------------|-------------|------------|------------|------------|-------|
|    | ~~~~~      |             |            | ~~~~~      |            | ~~~~~ |
| 1  | TCTAGAGCAT | CGGTAGGAGA  | AAATAAATG  | AAACAAGCA  | CTATTGCACT |       |
|    | AGATCTCGTA | CGCATCCCTCT | TTTATTTTAC | TTTGTTCGT  | GATAACGTGA |       |
| 51 | GGCACTCTTA | CCGTTGCTCT  | TCACCCCTGT | TACCAAGCC  | GAATTC     |       |
|    | CCGTGAGAAT | GGCAACGAGA  | AGTGGGACA  | ATGGTTTCGG | CTTAAG     |       |

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)



```

XbaI SphI

1 TCTAGAGCAT CCGTAGGAGA AAATAAATG AAACAAGCA CTATTGCACT
 AGATCTCGTA CGCATCCTCT TTTATTTAC TTTGTTTCGT GATAACGTGA

 Sapi

51 GGCACTCTTA CCGTTGCTCT TCACCCCTGT TACCAAAGCC GACTACAAAG
 CCGTGAGAAT GGCAACGAGA AGTGGGGACA ATGGTTTCGG CTGATGTTTC

 Muni EcoRI

101 ATGAAGTGCA ATTGGAATTC
 TACTTCACGT TAACCTTAAG

```

147 / 204

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

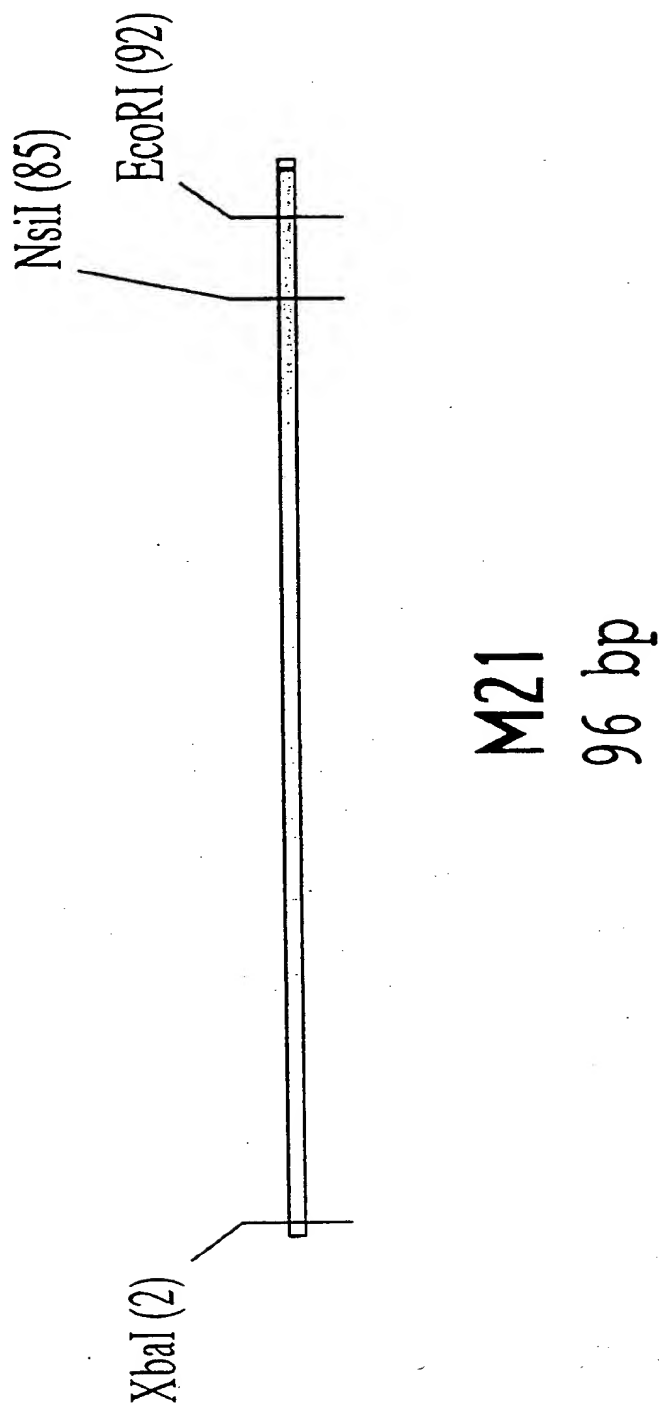




Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

M 21:

XbaI

-----

1 TCTAGAGGTT GAGGTGATTT TATGAAAAG AATATCGCAT TTCTTCTTGC  
AGATCTCCAA CTCCACTAAA ATACTTTTTC TTATAGCGTA AAGAAGAAGC

NsiI ECORI

-----

51 ATCTATGTTT GTTTTTCTA TTGCTACAAA TGCATACGCT GAATTC  
TAGATACAAG CAAAAGAGAT AACGATGTTT ACGTATGCCA CTTAAG

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

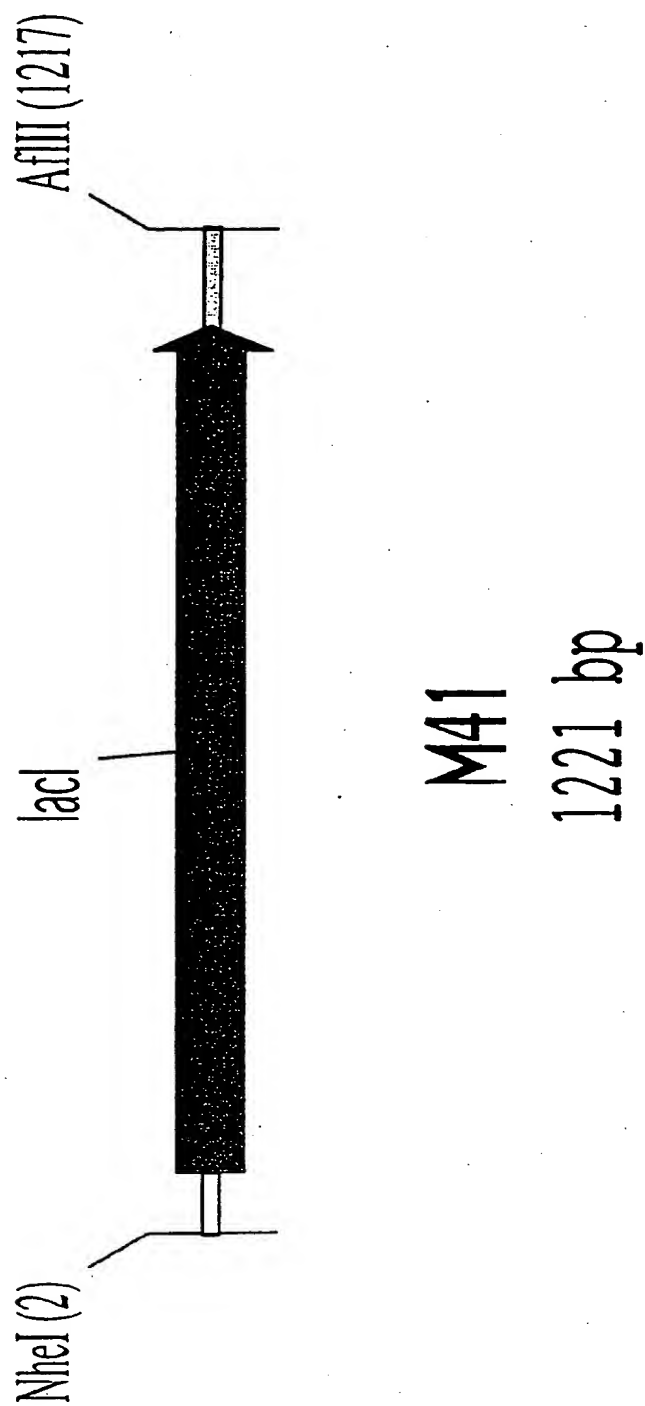


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|       |                                                        |                                                        |  |
|-------|--------------------------------------------------------|--------------------------------------------------------|--|
| M 41: |                                                        | NheI                                                   |  |
|       |                                                        | ~~~~~                                                  |  |
| 1     | GCTAGCATCG AATGGCGCAA AACCTTTCGC GGTATGGCAT GATAGCGCCC | CGATCGTAGC TTACCGCGTT TTGGAAAGCG CCATACCGTA CTATCGCGGG |  |
| 51    | GGAAGAGAGT CAATTCAGGG TGGTGAATGT GAAACCAGTA ACGTTATACG | CCTTCTCTCA GTTAAGTCCC ACCACTTACA CTTTGGTCAT TGCAATATGC |  |
| 101   | ATGTCGCAGA GTATGCCGGT GTCTCTTATC AGACCGTTTC CCGCGTGGTG | TACAGCGTCT CATACGGCCA CAGAGAATAG TCTGGCAAAG GCGCACCCAC |  |
| 151   | AACCAGGCCA GCCACGTTC TCGGAAAACG CGGAAAAAAG TGGAAGCGGC  | TTGGTCCGGT CGGTGCAAAG ACGCTTTTGC GCCCTTTTTC ACCTTCGCGC |  |
| 201   | GATGGCGGAG CTGAATTACA TTCCTAACCG CGTGGCACAA CAACTGGCGG | CTACCGCCTC GACTTAATGT AAGGATTGGC GCACCGTGTT GTTGACCGCC |  |
| 251   | GCAAACAGTC GTTGCTGATT GGCGTTGCCA CCTCCAGTCT GGCCCTGCAC | CGTTTGTCAG CAACGACTAA CCGCAACGGT GGAGGTCAGA CCGGGACGTG |  |
| 301   | GCGCCGTCGC AAATTGTCGC GCGGATTAAA TCTCGCGCCG ATCAACTGGG | CGGGCAGCG TTTAACAGCG CCGCTAATT AGAGCGCGGC TAGTTGACCC   |  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |             |             |            |             |             |
|-----|-------------|-------------|------------|-------------|-------------|
| 351 | TGCCAGCGTG  | GTCGTGTCGA  | TGGTAGAACG | AAGCGGCGTC  | GAAGCCTGTA  |
|     | ACGGTCGCAC  | CAGCACAGCT  | ACCATCTTGC | TTCGCCCGCAG | CTTCGGACAT  |
| 401 | AAGCGGCGGT  | GCACAAATCTT | CTCGCGCAAC | GTGTCAGTGG  | GCTGATTATT  |
|     | TTCGCCCGCCA | CGTGTTAGAA  | GAGCGCGTTG | CACAGTCACC  | CGACTAATAA  |
| 451 | AACTATCCGC  | TGGATGACCA  | GGATGCTATT | GCTGTGGAAG  | CTGCCCTGCAC |
|     | TTGATAGGCG  | ACCTACTGGT  | CCTACGATAA | CGACACCTTC  | GACGGACGTG  |
| 501 | TAATGTTCCG  | GCGTTATTTC  | TTGATGTCTC | TGACCAGACA  | CCCATCAACA  |
|     | ATTACAAGGC  | CGCAATAAAG  | AACTACAGAG | ACTGGTCTGT  | GGTAGTTGT   |
| 551 | GTATTATTTT  | CTCCCATGAG  | GACGGTACGC | GACTGGGCGT  | GGAGCATCTG  |
|     | CATAATAAAA  | GAGGGTACTC  | CTGCCATGCG | CTGACCCGCA  | CCTCGTAGAC  |
| 601 | GTCGCATTGG  | GCCACCAGCA  | AATCGCGCTG | TTAGCTGGCC  | CATTAAAGTTC |
|     | CAGCGTAACC  | CGGTGGTCGT  | TTAGCGCGAC | AATCGACCGG  | GTAATTCAAG  |
| 651 | TGTCTCGGCG  | CGTCTGCGTC  | TGGCTGGCTG | GCATAAATAT  | CTCACTCGCA  |
|     | ACAGAGCCCG  | GCAGACGCAG  | ACCGACCGAC | CGTATTTATA  | GAGTGAGCGT  |
| 701 | ATCAAATTCA  | GCCGATAGCG  | GAACGGGAAG | GCGACTGGAG  | TGCCATGTCC  |
|     | TAGTTTAAGT  | CGGCTATCGC  | CTTGCCCTTC | CGCTGACCTC  | ACGGTACAGG  |

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |            |             |             |             |             |
|------|------------|-------------|-------------|-------------|-------------|
| 751  | GGTTTCAAC  | AAACCATGCA  | AATGCTGAAT  | GAGGGCATCG  | TTCCCCACTGC |
|      | CCAAAAGTTG | TTTGGTACGT  | TTACGACTTA  | CTCCCCGTAGC | AAGGGTGACG  |
| 801  | GATGCTGGTT | GCCAACGATC  | AGATGGCGCT  | GGGCGCAATG  | CGTGCCATTA  |
|      | CTACGACCAA | CGGTGCTAG   | TCTACCGCGA  | CCCGCGTTAC  | GCACGGTAAT  |
| 851  | CCGAGTCCGG | GCTGCGCGTT  | GGTGCGGACA  | TCTCGGTAGT  | GGGATACGAC  |
|      | GGCTCAGGCC | CGACGCGCAA  | CCACGCCCTGT | AGAGCCATCA  | CCCTATGCTG  |
| 901  | GATACCGAGG | ACAGTCAATG  | TTATATCCCG  | CCGCTGACCA  | CCATCAAACA  |
|      | CTATGGCTCC | TGTCGAGTAC  | AATATAGGC   | GGCGACTGGT  | GGTAGTTTGT  |
| 951  | GGATTTTCGC | CTGCTGGGGC  | AAACCAGCGT  | GGACCGCTTG  | CTGCAACTCT  |
|      | CCTAAAAGCG | GACGACCCCG  | TTTGGTCGCA  | CCTGGCGAAC  | GACGTTGAGA  |
| 1001 | CTCAGGGCCA | GGCGGTGAAG  | GGCAATCAGC  | TGTTGCCCGT  | CTCACTGGTG  |
|      | GAGTCCCGGT | CCGCCACTTC  | CCGTAGTCG   | ACAACGGGCA  | GAGTGACCAC  |
| 1051 | AAAAGAAAAA | CCACCCCTGGC | TCCCAATACG  | CAAACCGCCT  | CTCCCCCGCG  |
|      | TTTTCTTTTT | GGTGGGACCG  | AGGTTATGC   | GTTTGGCGGA  | GAGGGGCGCG  |
| 1101 | GTTGGCCGAT | TCACTGATGC  | AGCTGGCAGC  | ACAGGTTTCC  | CGACTGGAAA  |
|      | CAACCGGCTA | AGTGACTACG  | TCGACCGTGC  | TGTCCAAAGG  | GCTGACCTTT  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

1151 GCGGGCAGTG AGGCTACCCG ATAAAGCGG CTTCCTGACA GGAGGCCGTT  
CGCCCGTCAC TCCGATGGC TATTTTCGCC GAAGGACTGT CCTCCGGCAA

AflII

~~~~~

1201 TTGTTTGTGCA GCCCACTTAA G  
AACAAAACGT CGGTGAATT C

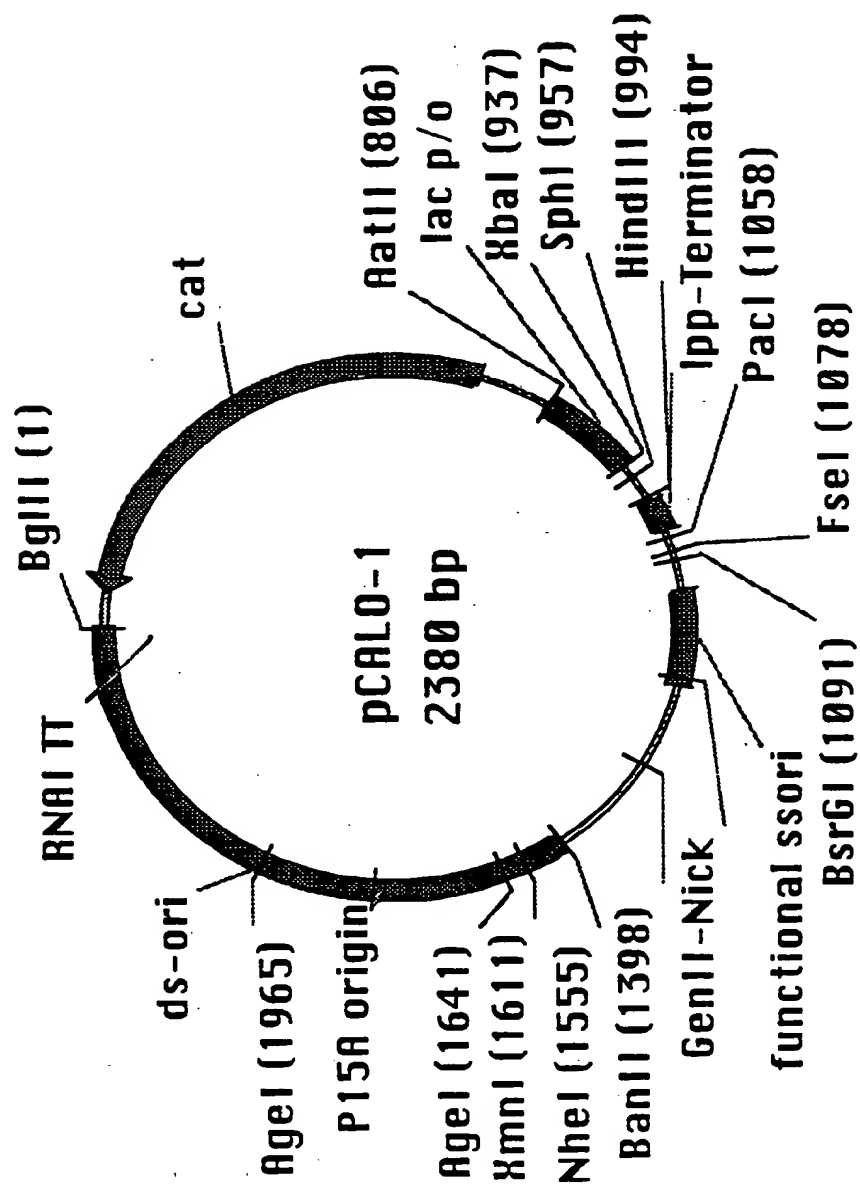


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|          |                                                                                                                     |
|----------|---------------------------------------------------------------------------------------------------------------------|
| pCALO-1: |                                                                                                                     |
| Bg1II    |                                                                                                                     |
| ~~~~~    |                                                                                                                     |
| 1        | GATCTAGCAC CAGGCGTTTA AGGGACCCAA TAACTGCCTT AAAAATAATTA<br>CTAGATCGTG GTCCGCAAAAT TCCCGTGGTT ATTGACGGAA TTTTTTTAAAT |
| 51       | CGCCCCGCCC TGCCACTCAT CGCAGTACTG TTGTAATTCA TTAAGCATTC<br>GCGGGGCGGG ACGGTGAGTA GCGTCATGAC AACATTAACT AATTTCGTAAG   |
| 101      | TGCCGACATG GAAGCCATCA CAAACGGCAT GATGAACCTG AATCGCCAGC<br>ACGGCTGTAC CTTCCGTTAGT GTTTGCCCGTA CTACTTGGAC TTAGCGGTTCG |
| 151      | GGCATCAGCA CCTTGTCGCC TTGCGTATAA TATTTGCCCA TAGTGAAAC<br>CCGTAGTCGT GGAACAGCGG AACGCATATT ATAAACGGGT ATCACTTTTG     |
| 201      | GGGGGCGAAG AAGTTGTCCA TATTGGCTAC GTTTAAATCA AAACCTGGTGA<br>CCCCCGCTTC TTCAACAGGT ATAACCGATG CAAATTAGT TTTGACCACT    |
| 251      | AACTCACCCA GGGATTGGCT GAGACGAAA ACATATTCTC AATAAACCCCT<br>TTGAGTGGGT CCTTAACCGA CTCTGCTTTT TGTATAAGAG TTATTTGGGA    |
| 301      | TTAGGGAAAT AGGCCAGGTT TTCACCGTAA CACGCCACAT CTGCGAATA<br>AATCCCTTTA TCCGGTCCAA AAGTGGCATT GTGCGGTGTA GAACGCTTAT     |



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |             |             |              |             |             |
|-----|-------------|-------------|--------------|-------------|-------------|
| 351 | TATGTGTAGA  | AACTGCCGGA  | AATCGTCGTG   | GTATTCACTC  | CAGAGCGATG  |
|     | ATACACATCT  | TTGACGGCCT  | TTAGCAGCAC   | CATAAGTGAG  | GTCTCGCTAC  |
| 401 | AAAACGTTTC  | AGTTTGCTCA  | TGGA AAAACGG | TGTAACAAGG  | GTGAACACTA  |
|     | TTTTTGCAAAG | TCAAACGAGT  | ACCTTTTGCC   | ACATTGTTCC  | CACTTGTGAT  |
| 451 | TCCCATATCA  | CCAGCTCACC  | GTCTTTCATT   | GCCATACGGA  | ACTCCGGGTG  |
|     | AGGTATAGT   | GGTCGAGTGG  | CAGAAAGTAA   | CGGTATGCCT  | TGAGGCCCCAC |
| 501 | AGCATTCATC  | AGCGGGGCAA  | GAATGTGAAT   | AAAGGCCGGA  | TAAAACTTGT  |
|     | TCGTAAGTAG  | TCCGCCCGTT  | CTTACACTTA   | TTTCCGGCCT  | ATTTTGAACA  |
| 551 | GCTTATTTTT  | CTTTACGGTC  | TTTAAAAAAGG  | CCGTAATATC  | CAGCTGAACG  |
|     | CGAATAAAAA  | GAAATGCCAG  | AAATTTTCC    | GGCATTTATAG | GTCGACTTGC  |
| 601 | GTCTGGTTAT  | AGGTACATTG  | AGCAACTGAC   | TGAAATGCCT  | CAAAATGTTT  |
|     | CAGACCAATA  | TCCATGTAAC  | TCGTTGACTG   | ACTTTACGGA  | GTTTACCAAG  |
| 651 | TTTACGATGC  | CATTGGGATA  | TATCAACGGT   | GGTATATCCA  | GTGATTTTTT  |
|     | AAATGCTACG  | GTAACCCCTAT | ATAGTTGCCA   | CCATATAGGT  | CACTAAAAAA  |
| 701 | TCTCCATTTT  | AGCTTCCTTA  | GCTCCTGAAA   | ATCTCGATAA  | CTCAAAAAAT  |
|     | AGAGGTAAAA  | TCGAAGGAAT  | CGAGGACTTT   | TAGAGCTATT  | GAGTTTTTTA  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |            |             |            |            |            |
|------|------------|-------------|------------|------------|------------|
| 751  | ACGCCCGGTA | GTGATCTTAT  | TTCATTATGG | TGAAAGTTGG | AACCTCACCC |
|      | TGCGGGCCAT | CACTAGAATA  | AAGTAATACC | ACTTTCACCC | TTGGAGTGGG |
|      | AatII      |             |            |            |            |
|      | ~~~~~      |             |            |            |            |
| 801  | GACGTCTAAT | GTGAGTTAGC  | TCACTCATTA | GGCACCCAG  | GCTTTACACT |
|      | CTGCAGATTA | CACTCAATCG  | AGTGAGTAAT | CCGTGGGGTC | CGAAATGTGA |
| 851  | TTATGCTTCC | GGCTCGTATG  | TTGTGTGGAA | TTGTGAGCGG | ATAACAATT  |
|      | AATACGAAGG | CCGAGCATAC  | AACACACCTT | AACACTCGCC | TATTGTTAAA |
|      | XbaI       |             |            |            |            |
|      | ~~~~~      |             |            |            |            |
| 901  | CACACAGGAA | ACAGCTATGA  | CCATGATTAC | GAATTTCTAG | ACCCCCCCCC |
|      | GTGTGTCCTT | TGTCGATACT  | GGTACTAATG | CTTAAAGATC | TGGGGGGGGG |
|      | SphI       |             |            |            |            |
|      | ~~~~~      |             |            |            |            |
| 951  | CGCATGCCAT | AAC TTCGTAT | AATGTACGCT | ATACGAAGTT | ATAAGCTTGA |
|      | GCGTACGGTA | TTGAAGCATA  | TTACATGCGA | TATGCTTCAA | TATTCGAAC  |
| 1001 | CCTGTGAAGT | GAAAAATGGC  | GCAGATTGTG | CGACATTTT  | TTTGTCTGCC |
|      | GGACACTTCA | CTTTTACC    | CGTCTAACAC | GCTGTAAAAA | AAACAGACGG |
|      | HindIII    |             |            |            |            |
|      | ~~~~~      |             |            |            |            |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      | PacI                                                                                                              | FseI  | BsrGI |
|------|-------------------------------------------------------------------------------------------------------------------|-------|-------|
|      | ~~~~~                                                                                                             | ~~~~~ | ~~~~~ |
| 1051 | GTTTAATTAA AGGGGGGGG GGGCCGGCCT GGGGGGGGT GTACATGAAA<br>CAAATTAAAT TCCCCCCCCC CCGGCCGGA CCCCCCCCCA CATGTACTTT     |       |       |
| 1101 | TTGTAAACGT TAATATTTG TTAAATTCG CGTTAAATTT TTGTAAATC<br>AACATTTGCA ATTATAAAC AATTTAAGC GCAATTTAAA AACAAATTTAG      |       |       |
| 1151 | AGCTCATTTT TTAACCAATA GGCCGAAATC GGCAAAATCC CTTATAAATC<br>TCGAGTAAAA AATTGGTTAT CCGCCTTAG CCGTTTAGG GAATATTTAG    |       |       |
| 1201 | AAAAGAATAG ACCGAGATAG GGTGAGTGT TGTTCCAGTT TGGAAACAAGA<br>TTTTCTTATC TGGCTCTATC CCAACTCACA ACAAGTCAA ACCTTGTTCT   |       |       |
| 1251 | GTCCACTATT AAAGAACGTG GACTCCAACG TCAAAGGCG AAAAACCGTC<br>CAGGTGATAA TTTCTTGAC CTGAGGTGC AGTTTCCCGC TTTTGGCAG      |       |       |
| 1301 | TATCAGGGCG ATGGCCCACT ACGAGAACCA TCACCCCTAAT CAAGTTTTTT<br>ATAGTCCCCG TACCCGGTGA TGCTCTTGGT AGTGGGATTA GTTCAAAAAA |       |       |
|      |                                                                                                                   |       | BanII |
|      |                                                                                                                   |       | ~~~~~ |
| 1351 | GGGGTCGAGG TGCCGTAAAG CACTAAATCG GAACCCCTAAA GGGAGCCCCC<br>CCCCAGCTCC ACGGCATTTC GTGATTTAGC CTGGGGATT CCCTCGGGGG  |       |       |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |                                                                                                                   |
|------|-------------------------------------------------------------------------------------------------------------------|
| 1401 | GATTAGAGC TTGACGGGGA AAGCCGGCGA ACGTGGCGAG AAAGGAAGGG<br>CTAAATCTCG AACTGCCCCCT TTCGGCCGCT TGCACCGCTC TTTCCTTCCC  |
| 1451 | AAGAAAGCGA AAGAGCGGG CGCTAGGGCG CTGGCAAGTG TAGCGGTCAC<br>TTCTTTTCGCT TTCCCTCGCC GCGATCCCCG GACCGTTCAC ATCGCCAGTG  |
| 1501 | GCTGCGCGTA ACCACCACAC CCGCCGCGCT TAATGCGCCG CTACAGGGCG<br>CGACGCGCAT TGGTGGTGTG GCGGCGCGGA ATTACGCGGC GATGTCCCCG  |
|      | NheI<br>~~~~~                                                                                                     |
| 1551 | CGTGCTAGCG GAGTGATAC TGGCTTACTA TGTGGCACT GATGAGGGTG<br>GCACGATCGC CTCACATATG ACCGAATGAT ACAACCGTGA CTACTCCCCAC   |
|      | XmnI<br>~~~~~                                                                                                     |
| 1601 | TCAGTGAAGT GCTTCATGTG GCAGGAGAAA AAAGGCTGCA CCGGTGCGTC<br>AGTCACTTCA CGAAGTACAC CGTCCTCTTT TTTCCGACGT GGCCACGCAG  |
| 1651 | AGCAGAATAT GTGATACAGG ATATATTCCG CTTCCCTCGCT CACTGACTCG<br>TCGTCTTATA CACTATGTCC TATATAAGGC GAAGGAGCGA GTGACTGAGC |
| 1701 | CTACGCTCGG TCGTTCGACT GCGGCGAGCG GAAATGGCTT ACGAACGGGG                                                            |
|      | AgeI<br>~~~~~                                                                                                     |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |               |              |            |            |             |
|------|---------------|--------------|------------|------------|-------------|
|      | GATGCGAGCC    | AGCAAGCTGA   | CGCCGCTCGC | CTTTACCGAA | TGCTTGCCCC  |
| 1751 | CGGAGATTTC    | CTGGAAGATG   | CCAGGAAGAT | ACTTAACAGG | GAAGTGAGAG  |
|      | GCCCTCTAAAG   | GACCTTCTAC   | GGTCCTTCTA | TGAATTGTCC | CTTCACTCTC  |
| 1801 | GGCCGCGGCA    | AAGCCGTTT    | TCCATAGGCT | CCGCCCCCCT | GACAAGCATC  |
|      | CCGGCGCCCGT   | TTCGGCAAA    | AGTATCCGA  | GGCGGGGGA  | CTGTTCCGTAG |
| 1851 | ACGAAATCTG    | ACGCTCAAAT   | CAGTGGTGGC | GAAACCCGAC | AGGACTATAA  |
|      | TGCTTTAGAC    | TGCGAGTTA    | GTCACCACCG | CTTTGGGCTG | TCCTGATATT  |
| 1901 | AGATACCAGG    | CGTTTCCCC    | TGGCGGCTCC | CTCCTGCGCT | CTCCTGTTCC  |
|      | TCTATGGTCC    | GCAAAGGGG    | ACCGCCGAGG | GAGACGCGA  | GAGACAAGG   |
|      | AgeI<br>~~~~~ |              |            |            |             |
| 1951 | TGCCTTTCGG    | TTTACC GG TG | TCATTCCGCT | GTTATGGCCG | CGTTTGCTCTC |
|      | ACGGAAGCC     | AAATGGCCAC   | AGTAAGGCCA | CAATACCGGC | GCAACACAGAG |
| 2001 | ATTCCACGCC    | TGACACTCAG   | TTCCGGGTAG | GCAGTTCGCT | CCAAGCTGGA  |
|      | TAAGGTGCGG    | ACTGTAGTC    | AAGCCCCATC | CGTCAAGCGA | GGTTCGACCT  |
| 2051 | CTGTATGCAC    | GAACCCCCCG   | TTCAGTCCGA | CCGCTGCGCC | TTATCCGGTA  |
|      | GACATACGTG    | CTTGGGGGGC   | AAGTCAGGCT | GGCGACGCGG | AATAGGCCAT  |

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|       |            |            |            |            |             |
|-------|------------|------------|------------|------------|-------------|
| 2101  | ACTATCGTCT | TGAGTCCAAC | CCGAAAGAC  | ATGCAAAAGC | ACCACTGGCA  |
|       | TGATAGCAGA | ACTCAGGTG  | GGCCTTTCTG | TACGTTTTCG | TGGTGACCCGT |
| 2151  | GCAGCCACTG | GTAATTGATT | TAGAGGAGTT | AGTCTTGAAG | TCATGCGCCG  |
|       | CGTCGGTGAC | CATTAACTAA | ATCTCCTCAA | TCAGAACTTC | AGTACGCGGC  |
| 2201  | GTTAAGGCTA | AACTGAAAGG | ACAAGTTTAA | GTGACTGCCG | TCCTCCAAGC  |
|       | CAATTCCGAT | TTGACTTTCC | TGTTCAAAAT | CACTGACGCG | AGGAGGTTCCG |
| 2251  | CAGTTACCTC | GGTTCAAAGA | GTTGGTAGCT | CAGAGAACCT | ACGAAAACC   |
|       | GTCAATGGAG | CCAAGTTTCT | CAACCATCGA | GTCTCTTGA  | TGCTTTTTCG  |
| 2301  | GCCCTGCAAG | GCGGTTTTTT | CGTTTTCAGA | GCAAGAGATT | ACGCGCAGAC  |
|       | CGGGACGTTT | CGCCAAAAAA | GCAAAAGTCT | CGTTCTCTAA | TGCGCGTCTG  |
| BglII |            |            |            |            |             |
| 2351  | CAAAACGATC | TCAAGAAGAT | CATCTTATTA |            |             |
|       | GTTTTCCTAG | AGTTCTTCTA | GTAGAATAAT |            |             |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

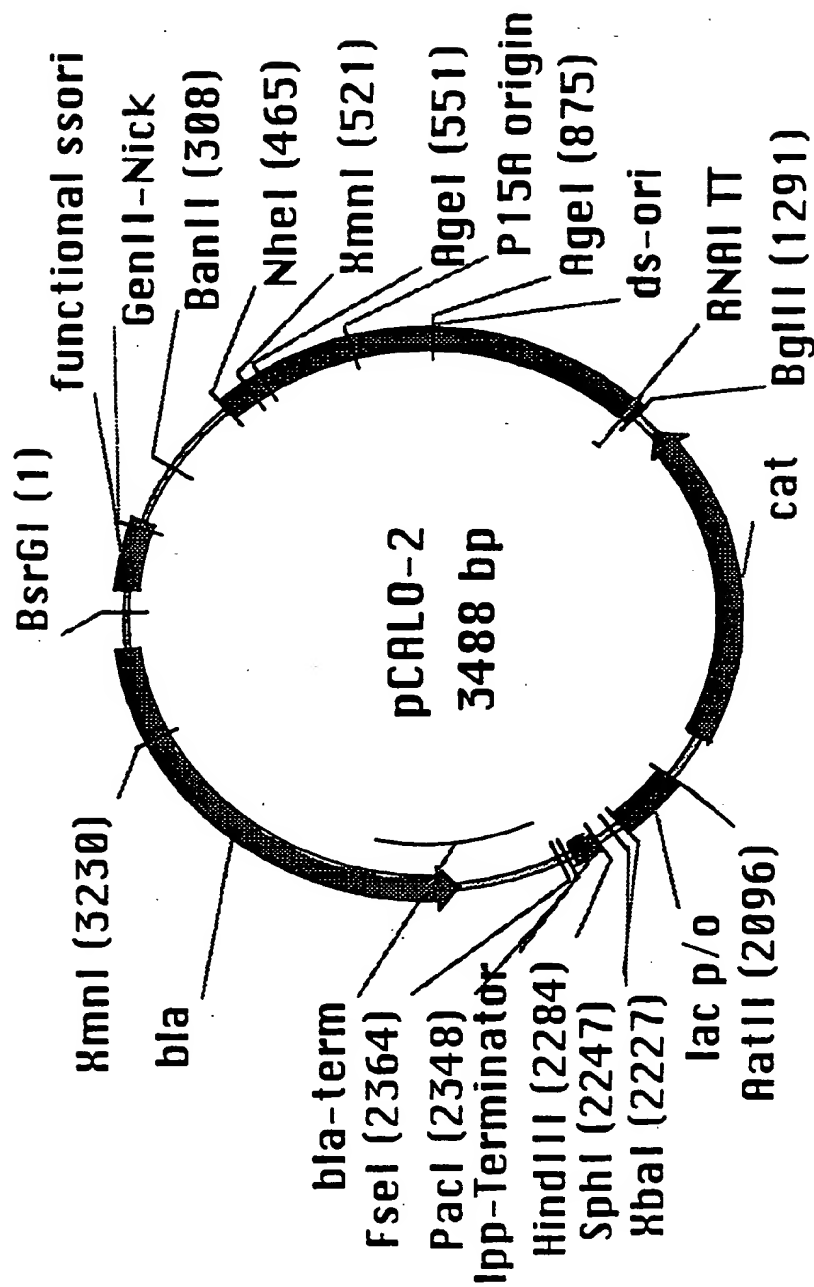


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

pCALO-2:

BsrGI

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1 GTACATGAAA TTGTAAACGT TAATATTTTG TTAAAATTCC CGTTAAATTT  
CATGTAATTT AACATTTGCA ATTATAAAC AATTTTAAGC GCAATTTAAA

51 TTGTTAAATC AGCTCATTTT TTAACCAATA GGCCGAAATC GGCAAAATCC  
AACAAATTTAG TCGAGTAAA AATTGGTTAT CCGGCTTTAG CCGTTTTAGG

101 CTTATAAATC AAAAGAATAG ACCGAGATAG GGTGAGTGT TGTTCCAGTT  
GAATATTTAG TTTTCTTATC TGGCTCTATC CCAACTCACA ACAAGGTCAA

151 TGGAAACAAGA GTCCACTATT AAAGAACGTG GACTCCAACG TCAAAGGGCG  
ACCTTGTTCT CAGGTGATAA TTCTTTGCAC CTGAGGTTGC AGTTTCCCGC

201 AAAAACCGTC TATCAGGGCG ATGGCCCACT ACGAGAACCA TCACCCTAAT  
TTTTTTGGCAG ATAGTCCCGC TACCGGGTGA TGCTCTTGGT AGTGGGATTA

251 CAAGTTTTTT GGGTCGAGG TGCCGTAAAG CACTAAATCG GAACCCATAA  
GTTCAAAAAA CCCAGCTCC ACGCATTTT GTGATTTAGC CTGCGGATTT

BanII

~~~~~

301 GGGAGCCCC GATTAGAGC TTGACGGGA AAGCCGGCA ACGTGGCGAG



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |             |             |             |            |            |
|-----|-------------|-------------|-------------|------------|------------|
|     | CCCTCGGGG   | CTAATCTCG   | AACTGCCCT   | TTCGGCGCT  | TGCACCGCTC |
| 351 | AAAGGAAGG   | AAGAAAGCGA  | AAGGAGCGG   | CGCTAGGGCG | CTGGCAAGTG |
|     | TTTCCCTTCCC | TTCTTTTCGCT | TTCCCTCGCCC | GCGATCCCGC | GACCGTTCAC |
| 401 | TAGCGGTCAC  | GCTGCGCGTA  | ACCACCACAC  | CCGCCGCGCT | TAATGCGCCG |
|     | ATCGCCAGTG  | CGACGCGCAT  | TGGTGGTGTG  | GGCGCGCGCA | ATTACGCGGC |
|     |             | NheI        |             |            |            |
|     |             | ~~~~~       |             |            |            |
| 451 | CTACAGGGCG  | CGTGCTAGCG  | GAGTGTATAC  | TGGCTTACTA | TGTTGGCACT |
|     | GATGTCCCCG  | GCACGATCGC  | CTCACATATG  | ACCGAATGAT | ACAACCGTGA |
|     |             | XmnI        |             |            | AgeI       |
|     |             | ~~~~~       |             |            | ~          |
| 501 | GATGAGGGTG  | TCAGTGAAGT  | GCTTCATGTG  | GCAGGAGAAA | AAAGGCTGCA |
|     | CTACTCCAC   | AGTCACTTCA  | CGAAGTACAC  | CGTCCCTCTT | TTTCCGACGT |
|     | AgeI        |             |             |            |            |
|     | ~~~~~       |             |             |            |            |
| 551 | CCGGTGGGTC  | AGCAGAATAT  | GTGATACAGG  | ATATATTCCG | CTTCCTCGCT |
|     | GGCCACGCAG  | TCGTCTTATA  | CACTATGTCC  | TATATAAGGC | GAAGGAGCGA |
| 601 | CACTGACTCG  | CTACGCTCGG  | TCGTTGACT   | GCGCGAGCG  | GAAATGGCTT |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |               |            |            |             |            |
|-----|---------------|------------|------------|-------------|------------|
|     | GTGACTGAGC    | GATGCGAGCC | AGCAAGCTGA | CGCCGCTCGC  | CTTACCGAA  |
| 651 | ACGAACGGG     | CGGAGATTTC | CTGGAAGATG | CCAGGAAGAT  | ACTTAACAGG |
|     | TGCTTGCCCC    | GCCTCTAAG  | GACCTTCTAC | GGTCCCTTCTA | TGAATTGTCC |
| 701 | GAAGTGAGAG    | GGCCGCGGCA | AAGCCGTTT  | TCCATAGGCT  | CCGCCCCCCT |
|     | CTTCACTCTC    | CCGGCGCCGT | TTCGGCAAAA | AGGTATCCGA  | GGCGGGGGA  |
| 751 | GACAAGCATC    | ACGAAATCTG | ACGCTCAAAT | CAGTGGTGGC  | GAAACCCGAC |
|     | CTGTTCTGTAG   | TGCTTTAGAC | TGCGAGTTTA | GTCACCAACG  | CTTTGGGCTG |
| 801 | AGGACTATAA    | AGATACCAGG | CGTTTCCCCC | TGGCGGCTCC  | CTCCTGCGCT |
|     | TCCTGATATT    | TCTATGGTCC | GCAAAGGGGG | ACCGCCGAGG  | GAGGACGCGA |
|     | AgeI<br>~~~~~ |            |            |             |            |
| 851 | CTCCTGTTC     | TGCCTTTCGG | TTTACCGGTG | TCATTCCGCT  | GTTATGGCCG |
|     | GAGGACAAG     | ACGGAAAGCC | AAATGGCCAC | AGTAAGGCCA  | CAATACCGGC |
| 901 | CGTTTGTCTC    | ATTCCACGCC | TGACACTCAG | TTCCGGGTAG  | GCAGTTCGCT |
|     | GCAAACAGAG    | TAAGGTGCGG | ACTGTAGTC  | AAGGCCCATC  | CGTCAAGCGA |
| 951 | CCAAGCTGGA    | CTGTATGCAC | GAACCCCCCG | TTCAGTCCGA  | CCGCTGCGCC |
|     | GGTTCGACCT    | GACATACGTG | CTTGGGGGGC | AAGTCAGGCT  | GGCGACGCGG |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |             |             |             |             |             |
|------|-------------|-------------|-------------|-------------|-------------|
| 1001 | TTATCCGGTA  | ACTATCGTCT  | TGAGTCCAAC  | CCGGAAGAC   | ATGCAAAAGC  |
|      | AATAGGCCAT  | TGATAGCAGA  | ACTCAGGTG   | GGCCTTTCTG  | TACGTTTTTCG |
| 1051 | ACCACTGGCA  | GCAGCCACTG  | GTAATTGATT  | TAGAGGAGTT  | AGTCTTGAAG  |
|      | TGGTGACCGT  | CGTCGGTGAC  | CATTAACTAA  | ATCTCCTCAA  | TCAGAACTTC  |
| 1101 | TCATGCGCCG  | GTTAAGGCTA  | AACTGAAAGG  | ACAAGTTTTA  | GTGACTGCGC  |
|      | AGTACGCGGC  | CAATTCCGAT  | TTGACTTTCC  | TGTTCAAAAT  | CACTGACGCG  |
| 1151 | TCCTCCAAGC  | CAGTTACCTC  | GGTCAAAGA   | GTTGGTAGCT  | CAGAGAACCT  |
|      | AGGAGGTTTC  | GTC AATGGAG | CCAAGTTTCT  | CAACCATCGA  | GTCCTCTTGA  |
| 1201 | ACGAAAACC   | GCCCTGCAAG  | GCGGTTTTTT  | CGTTTTTCTA  | GCAAGAGATT  |
|      | TGCTTTTTTG  | CGGACGTTTC  | CGCCAAAATA  | GCAAAAGTCT  | CGTTCCTCTAA |
|      |             |             |             | BglII       |             |
|      |             |             |             | ~~~~~       |             |
| 1251 | ACGCGCAGAC  | CAAACGATC   | TCAAGAAGAT  | CATCTTATTA  | GATCTAGCAC  |
|      | TGCGCGTCTG  | GTTTTGCTAG  | AGTTCTTCTA  | GTAGAATAAT  | CTAGATCGTG  |
| 1301 | CAGGCGTTTA  | AGGCAACCAA  | TAACTGCCCTT | AAAAAATAA   | CGCCCCGCCC  |
|      | GTCCGCAAAAT | TCCCGTGGTT  | ATTGACGGAA  | TTTTTTTAAAT | GCGGGGCGGG  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |              |             |             |             |             |
|------|--------------|-------------|-------------|-------------|-------------|
| 1351 | TGCCACTCAT   | CGCAGTACTG  | TTGTAATTCA  | TTAAGCATTC  | TGCCGACATG  |
|      | ACGGTGAGTA   | GGTCATGAC   | AACATTAAGT  | AATTCGTAAG  | ACGGCTGTAC  |
| 1401 | GAAGCCATCA   | CAAACGGCAT  | GATGAACCTG  | AATCGCCAGC  | GGCATCAGCA  |
|      | CTTCGGTAGT   | GTTTGCCGTA  | CTACTTGGAC  | TTAGCGGTGG  | CCGTAGTCGT  |
| 1451 | CCTTGTCGCC   | TTGCGTATAA  | TATTTGCCCA  | TAGTGAAAC   | GGGGCGGAAG  |
|      | GGAACAGCGG   | AACGCATATT  | ATAAACGGGT  | ATCACTTTTG  | CCCCCGCTTC  |
| 1501 | AAGTTGTCCA   | TATTGGCTAC  | GTTTAAATCA  | AAACTGGTGA  | AACTCACCCA  |
|      | TTCAACACAGGT | ATAACCGATG  | CAAATTTAGT  | TTTGACCACT  | TTGAGTGGGT  |
| 1551 | GGGATTGGCT   | GAGACGAAA   | ACATATTCTC  | AATAAACCCCT | TTAGGGAAAT  |
|      | CCCTAACCGA   | CTCTGCTTTT  | TGTATAAGAG  | TTATTTGGGA  | AATCCCCTTA  |
| 1601 | AGGCCAGGTT   | TTCAACCGTAA | CACGCCACAT  | CTTGCGAATA  | TATGTGTAGA  |
|      | TCCGGTCCAA   | AAGTGGCATT  | GTGCGGTGTA  | GAACGCTTAT  | ATACACATCT  |
| 1651 | AACTGCCCGGA  | AATCGTCGTG  | GTATTCACCTC | CAGAGCGATG  | AAAACGTTTC  |
|      | TTGACGGCCT   | TTAGCAGCAC  | CATAAGTGAG  | GTCTCGCTAC  | TTTTTGCAAAG |
| 1701 | AGTTTGCTCA   | TGGAACACGG  | TGTAACAAGG  | GTGAACACTA  | TCCCATATCA  |
|      | TCAAACGAGT   | ACCTTTTGCC  | ACATTGTTCC  | CACCTGTGAT  | AGGGTATAGT  |

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |             |             |            |             |            |
|------|-------------|-------------|------------|-------------|------------|
| 1751 | CCAGCTCACC  | GTCCTTTCATT | GCCATACGGA | ACTCCGGGTG  | AGCATTCATC |
|      | GGTCGAGTGG  | CAGAAAGTAA  | CGGTATGCCT | TGAGGCCCCAC | TCGTAAGTAG |
| 1801 | AGGCGGGCAA  | GAATGTGAAT  | AAAGCCGGA  | TAAAACTTGT  | GCTTATTTT  |
|      | TCCGCCCGTT  | CTTACACTTA  | TTTCCGGCCT | ATTTGAACA   | CGAATAAAAA |
| 1851 | CTTTACGGTC  | TTTAAAAAGG  | CCGTAATATC | CAGCTGAACG  | GTCTGGTTAT |
|      | GAAATGCCAG  | AAATTTTCC   | GGCATTATAG | GTCGACTTGC  | CAGACCAATA |
| 1901 | AGGTACATTG  | AGCAACTGAC  | TGAAATGCCT | CAAAATGTTT  | TTTACGATGC |
|      | TCCATGTAAC  | TCGTTGACTG  | ACTTTACGGA | GTTTACAAAG  | AAATGCTACG |
| 1951 | CATTGGGATA  | TATCAACGGT  | GGTATATCCA | GTGATTTT    | TCTCCATTTT |
|      | GTAACCCCTAT | ATAGTTGCCA  | CCATATAGGT | CACTAAAAAA  | AGAGGTAAAA |
| 2001 | AGCTTCCTTA  | GCTCCTGAAA  | ATCTCGATAA | CTCAAAAAAT  | ACGCCCGGTA |
|      | TCGAAGGAAT  | CGAGGACTTT  | TAGAGCTATT | GAGTTTTT    | TGCGGGCCAT |
|      |             |             |            | AatII       |            |
|      |             |             |            | ~~~~~       |            |
| 2051 | GTGATCTTAT  | TTCAATTATGG | TGAAAGTTGG | AACCTCACCC  | GACGTCTAAT |
|      | CACTAGAATA  | AAGTAATACC  | ACTTTCAACC | TTGGAGTGGG  | CTGCAGATTA |
| 2101 | GTGAGTTAGC  | TCACTCATTA  | GGCACCCAG  | GCTTTACACT  | TTATGCTTCC |

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Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |                                                                                                                   |       |
|------|-------------------------------------------------------------------------------------------------------------------|-------|
|      | CACTCAATCG AGTGAGTAAT CCGTGGGGTC CGAAATGTGA AATACGAAGG                                                            |       |
| 2151 | GGCTCGTATG TTGTGTGGAA TTGTGAGCGG ATAACAATTT CACACAGGAA<br>CCGAGCATAAC AACACACCTT AACACTCGCC TATTGTTAAA GTGTGTCCTT |       |
|      | XbaI                                                                                                              | SphI  |
|      | ~~~~~                                                                                                             | ~~~~~ |
| 2201 | ACAGCTATGA CCATGATTAC GAATTCTAG ACCCCCCCCC CGCATGCCAT<br>TGTCGATACT GGTACTAATG CTAAAGATC TGGGGGGGGG GCGTACGGTA    |       |
|      | HindIII                                                                                                           |       |
|      | ~~~~~                                                                                                             |       |
| 2251 | AAC TTCGTAT AATGACGCT ATACGAAGTT ATAAGCTTGA CCTGTGAAGT<br>TTGAAGCATA TTACATGCCA TATGCTTCAA TATTCGAACT GGACACTTCA  |       |
|      |                                                                                                                   | PacI  |
|      |                                                                                                                   | ~~~~~ |
| 2301 | GAAAAATGGC GCAGATTGTG CGACATTTT TTTGTCTGCC GTTTAATTAA<br>CTTTTACCG CGTCTAACAC GCTGTAAAAA AACACAGACGG CAAATTAAAT   |       |
|      | FseI                                                                                                              |       |
|      | ~~~~~                                                                                                             |       |
| 2351 | GGGGGGGGGC CGGCCATTAT CAAAAAGGAT CTCAAGAAGA TCCTTTGATC<br>CCCCCCCCCG GCCGGTAATA GTTTTTCCTA GAGTTCTTCT AGGAAACTAG  |       |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |             |            |             |             |             |
|------|-------------|------------|-------------|-------------|-------------|
| 2401 | TTTTTCTACGG | GGTCTGACGC | TCAGTGGAAC  | GAAAACTCAC  | GTTAAGGGAT  |
|      | AAAAGATGCC  | CCAGACTGCG | AGTCACCTTG  | CTTTTGAGTG  | CAATTCCCCTA |
| 2451 | TTTGGTCAATG | AGATTATCAA | AAAGGATCTT  | CACCTAGATC  | CTTTTAAATT  |
|      | AAACCAGTAC  | TCTAATAGTT | TTTCCCTAGAA | GTGGATCTAG  | GAAAATTATA  |
| 2501 | AAAAATGAAG  | TTTTAAATCA | ATCTAAAGTA  | TATATGAGTA  | AACTTGGTCT  |
|      | TTTTTACTTC  | AAAATTAGT  | TAGATTTTCAT | ATATACTCAT  | TTGAACCAGA  |
| 2551 | GACAGTTACC  | CAATGCTTAA | TCAGTGAGGC  | ACCTATCTCA  | GCGATCTGTC  |
|      | CTGTCAATGG  | GTACGAATT  | AGTCACTCCG  | TGGATAGAGT  | CGCTAGACAG  |
| 2601 | TATTTTCGTTT | ATCCATAGTT | GCCTGACTCC  | CCGTCGTGTA  | GATAACTACG  |
|      | ATAAAGCAAG  | TAGGTATCAA | CGGACTGAGG  | GCGAGCACAT  | CTATTGATGC  |
| 2651 | ATACGGGAGG  | GCTTACCATC | TGGCCCCAGT  | GCTGCAATGA  | TACCCGCGAGA |
|      | TATGCCCTCC  | CGAATGGTAG | ACCGGGGTCA  | CGACGTTACT  | ATGGCGCTCT  |
| 2701 | CCCACGCTCA  | CCGGCTCCAG | ATTATCAGC   | AATAAACCCAG | CCAGCCGGAA  |
|      | GGGTGCGAGT  | GGCCGAGGTC | TAAATAGTCG  | TTATTGGTC   | GGTCGGCCTT  |
| 2751 | GGGCCGAGCG  | CAGAAGTGGT | CCTGCAACTT  | TATCCGCCCTC | CATCCAGTCT  |
|      | CCCCGCTCGC  | GTCTTCACCA | GGACGTTGAA  | ATAGCGGGAG  | GTAGGTCAGA  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |             |             |             |             |             |
|------|-------------|-------------|-------------|-------------|-------------|
| 2801 | ATTAACGTGTT | GCCGGGAAGC  | TAGAGTAAGT  | AGTTCGCCAG  | TTAATAGTTT  |
|      | TAATTGACAA  | CGGCCCTTCG  | ATCTCATTCA  | TCAAGCGGTC  | AATTATCAAA  |
| 2851 | GCGCAACGTT  | GTTGCCATTG  | CTACAGGCAT  | CGTGGTGTC   | CGCTCGTCGT  |
|      | CGCGTTGCAA  | CAACGGTAAC  | GATGTCCGTA  | GCACCACAGT  | GCGAGCAGCA  |
| 2901 | TTGGTATGGC  | TTCAATTCAGC | TCCGGTTCCC  | AACGATCAAG  | GCGAGTTACA  |
|      | AACCATAACG  | AAGTAAGTCG  | AGGCCAAGGG  | TTGCTAGTTC  | CGCTCAATGT  |
| 2951 | TGATCCCCCA  | TGTTGTGCAA  | AAAAGCGGTT  | AGTCTCCTTCG | GTCTCTCCGAT |
|      | ACTAGGGGGT  | ACAACACGTT  | TTTTTCGCCAA | TCGAGGAAGC  | CAGGAGGCTA  |
| 3001 | CGTTGTCAGA  | AGTAAGTTGG  | CCGCAGTGTT  | ATCACTCATG  | GTTATGGCAG  |
|      | GCAACAGTCT  | TCATTCAACC  | GGCGTCACAA  | TAGTGAGTAC  | CAATACCGTC  |
| 3051 | CACTGCATAA  | TTCTCTTACT  | GTCAATGCCAT | CCGTAAGATG  | CTTTTCTGTG  |
|      | GTGACGTATT  | AAGAGAAATGA | CAGTACGGTA  | GGCATTCTAC  | GAAAAGACAC  |
| 3101 | ACTGGTGAGT  | ACTCAACCAA  | GTCAATCTGA  | GAATAGTGA   | TGCGGCGACC  |
|      | TGACCACTCA  | TGAGTTGGTT  | CAGTAAGACT  | CTTATCACAT  | ACGCCGCTGG  |
| 3151 | GAGTTGCTCT  | TGCCCCGGCGT | CAATACGGGA  | TAATACCGCG  | CCACATAGCA  |
|      | CTCAACGAGA  | ACGGGCCGCA  | GTTATGCCCT  | ATTATGGCGC  | GGTGTATCGT  |



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

| XmnI  |                                                                                                                   |
|-------|-------------------------------------------------------------------------------------------------------------------|
| ~~~~~ |                                                                                                                   |
| 3201  | GAAC TT TAAA AGTGCTCATC ATTGAAAAC GTTCTTCGGG GCGAAACTC<br>CTTGAAATTT TCACGAGTAG TAACCTTTTG CAAGAAGCCC CGCTTTTGAG  |
| 3251  | TCAAGGATCT TACCGCTGTT GAGATCCAGT TCGATGTAAC CCACTCGCGC<br>AGTTCCCTAGA ATGGCGACAA CTC TAGGTCA AGCTACATG GTGAGCGCG  |
| 3301  | ACCCAAC TGA TCCTCAGCAT CTTT TACTTT CACCAGCGTT TCTGGGTGAG<br>TGGGTTGACT AGGAGTCGTA GAAATGAAA GTGGTCGCAA AGACCCACTC |
| 3351  | CAAAAACAGG AAGGCAAAAT GCCGCAAAA AGGGAATAAG GCGACACGG<br>GTTT TGTCC TTCCGTTT TA CGCGGTTTTT TCCCTTATTC CCGCTGTGCC   |
| 3401  | AAATGTTGAA TACTCATACT CTTCCCTTTT CAATATTATT GAAGCATTTA<br>TTTACAACCTT ATGAGTATGA GAAGGAAAAA GTTATAATAA CTCGTAAAT  |
| BsrGI |                                                                                                                   |
| 3451  | TCAGGGTTAT TGTCTCATGA GCGGATACAT ATTTGAAT<br>AGTCCCAATA ACAGAGTACT CGCCTATGTA TAAACTTA                            |

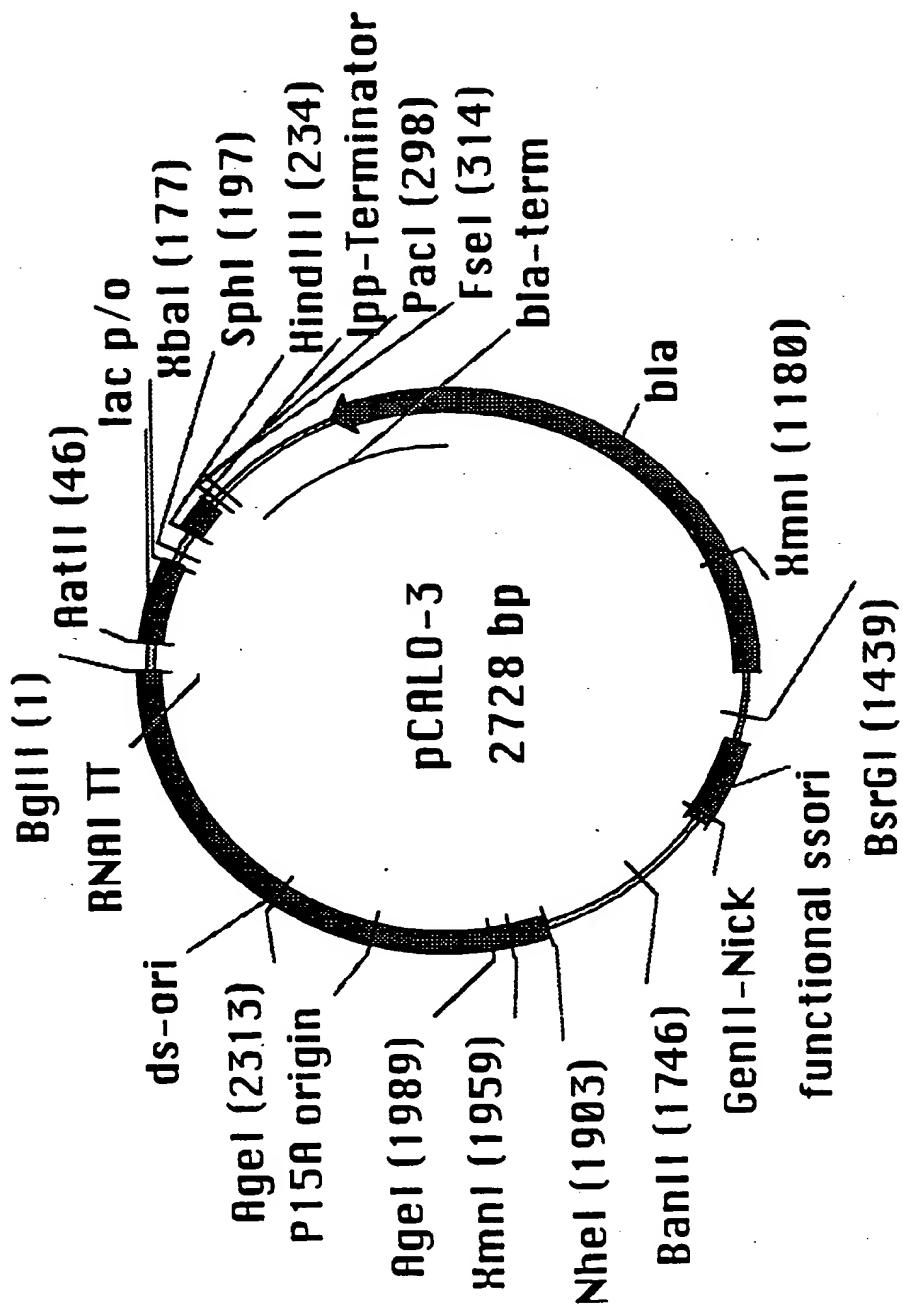


Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

| pCALO-3: |                                                          | AatII   |  |
|----------|----------------------------------------------------------|---------|--|
|          |                                                          | ~~~~~   |  |
|          |                                                          | BglII   |  |
|          |                                                          | ~~~~~   |  |
| 1        | GATCTCATAA CTTCTGTATAA TGTATGCTAT ACGAAGTTAT GACGTCTAAT  |         |  |
|          | CTAGAGTATT GAAGCATATT ACATACGATA TGCTTCAATA CTGCAGATTA   |         |  |
| 51       | GTGAGTTAGC TCACTCATTA GGCACCCAG GCTTTACACT TTATGCTTCC    |         |  |
|          | CACTCAATCG AGTGAGTAAT CCGTGGGTC CGAATGTGA AATACGAAGG     |         |  |
| 101      | GGCTCGTATG TTGTGTGGAA TTGTGAGCGG ATAACAATTT CACACAGGAA   |         |  |
|          | CCGAGCATAC AACACACCCCTT AACACTCGCC TATTGTTAAA GTGTGTCCTT |         |  |
|          |                                                          | SphI    |  |
|          |                                                          | ~~~~~   |  |
| 151      | ACAGCTATGA CCATGATTAC GAATTCTAG ACCCCCCCCC CGCATGCCAT    |         |  |
|          | TGTCGATACT GGTAATAATG CTAAAGATC TGGGGGGGGG GCGTACGGTA    |         |  |
|          |                                                          | HindIII |  |
|          |                                                          | ~~~~~   |  |
| 201      | AACTTCGTAT AATGTACGCT ATACGAAGTT ATAAGCTTGA CCTGTGAAGT   |         |  |
|          | TTGAAGCATA TTACATGCGA TATGCTTCAA TATTCGAACT GGACACTTCA   |         |  |
|          |                                                          | PacI    |  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     | FseI        |            |            |            |            |            |             |            |            |             |
|-----|-------------|------------|------------|------------|------------|------------|-------------|------------|------------|-------------|
|     | 1           | 2          | 3          | 4          | 5          | 6          | 7           | 8          | 9          | 10          |
| 251 | GAAAAATGGC  | GCAGATTGTG | CGACATTTT  | TTTGTCTGCC | GTTTAATTAA | CTTTTACCG  | CGCTAACAC   | GCTGTAAAA  | AAACAGACGG | CAAATTAATT  |
| 301 | GGGGGGGGC   | CGCCATTAT  | CAAAAAGGAT | CTCAAGAAGA | TCCTTTGATC | CCCCCCCCG  | GCCGGTAATA  | GTTTTCCTA  | GAGTTCTTCT | AGGAAACTAG  |
| 351 | TTTTTCTACGG | GGTCTGACGC | TCAGTGGAAC | GAAAACTCAC | GTTAAGGGAT | AAAAGATGCC | CCAGACTGCG  | AGTCACCTTG | CTTTTGAGTG | CAATTCCCCTA |
| 401 | TTTGGTCATG  | AGATTATCAA | AAAGGATCTT | CACCTAGATC | CTTTTAAATT | AAACCAGTAC | TCTAATAGTT  | TTTCCTAGAA | GTGGATCTAG | GAAAATTTAA  |
| 451 | AAAAATGAAG  | TTTTAAATCA | ATCTAAAGTA | TATATGAGTA | AACTTGGTCT | TTTTTACTTC | AAAATTTAGT  | TAGATTTCAT | ATATACTCAT | TTGAACCCAGA |
| 501 | GACAGTTACC  | CAATGCTTAA | TCAGTGAGGC | ACCTATCTCA | GCGATCTGTC | CTGTCAATGG | GTTACGGAATT | AGTCACTCCG | TGGATAGAGT | CGCTAGACAG  |
| 551 | TATTTTCGTTT | ATCCATAGTT | GCCTGACTCC | CCGTCGTGTA | GATAACTACG | ATAAAGCAAG | TAGGTATCAA  | CGGACTGAGG | GGCAGCACAT | CTATTGATGC  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|     |            |             |             |             |             |
|-----|------------|-------------|-------------|-------------|-------------|
| 601 | ATACGGGAGG | GCTTACCATC  | TGGCCCCCAGT | GCTGCAATGA  | TACCGCGAGA  |
|     | TATGCCCTCC | CGAATGGTAG  | ACCGGGGTCA  | CGACGTTACT  | ATGGCGCTCT  |
| 651 | CCCACGCTCA | CCGGCTCCAG  | ATTTATCAGC  | AATAAACCCAG | CCAGCCGGAA  |
|     | GGGTGCGAGT | GGCCGAGGTC  | TAAATAGTCG  | TTATTTGGTC  | GGTCGGCCCTT |
| 701 | GGGCCGAGCG | CAGAAGTGGT  | CCTGCAACTT  | TATCCGCCCTC | CATCCAGTCT  |
|     | CCCGGCTCGC | GTCCTCACCA  | GGACGTTGAA  | ATAGGCGGAG  | GTAGGTCAGA  |
| 751 | ATTAAGTGT  | GCCGGGAAGC  | TAGAGTAAGT  | AGTTCGCCAG  | TTAATAGTTT  |
|     | TAATTGACAA | CGGCCCTTCG  | ATCTCATTTCA | TCAAGCGGTC  | AATTATCAAA  |
| 801 | GCGCAACGTT | GTTGCCATTG  | CTACAGGCAT  | CGTGGTGTC   | CGCTCGTCGT  |
|     | CGCGTTGCAA | CAACGGTAAC  | GATGTCCGTA  | GCACCACAGT  | GCGAGCAGCA  |
| 851 | TTGGTATGGC | TTCATTTCAGC | TCCGGTTCCC  | AACGATCAAG  | GCGAGTTACA  |
|     | AACCATACCG | AAGTAAGTCG  | AGGCCAAGGG  | TTGCTAGTTC  | CGCTCAATGT  |
| 901 | TGATCCCCCA | TGTTGTGCAA  | AAAAGCGGTT  | AGCTCCTTCG  | GTCCCTCCGAT |
|     | ACTAGGGGGT | ACAACACGTT  | TTTTTCGCCAA | TCGAGGAAGC  | CAGGAGGCTA  |
| 951 | CGTTGTCAGA | AGTAAGTTGG  | CCGCAGTGTT  | ATCACTCATG  | GTTATGGCAG  |
|     | GCAACAGTCT | TCATTCAACC  | GGCGTCAACA  | TAGTGAGTAC  | CAATACCGTC  |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |                                                        |                                                        |
|------|--------------------------------------------------------|--------------------------------------------------------|
| 1001 | CACTGCATAA TTCTCTTACT GTCATGCCAT CCGTAAGATG CTTTCTGTG  | GTGACGTATT AAGAGAAATGA CAGTACGGTA GGCATTCTAC GAAAGACAC |
| 1051 | ACTGGTGAGT ACTCAACCAA GTCATTCTGA GAATAGTGA TGCGGCGACC  | TGACCACTCA TGAGTTGGTT CAGTAAGACT CTTATCACAT ACGCCGCTGG |
| 1101 | GAGTTGCTCT TGCCCGGCGT CAATACGGGA TAATACCGCG CCACATAGCA | CTCAACGAGA ACGGCGCGCA GTTATGCCCT ATTATGGCGG GGTGTATCGT |
| XmnI |                                                        |                                                        |
| 1151 | GAACTTTAAA AGTGCTCATC ATTGGAAAAC GTTCTTCGGG GCGAAACTC  | CTTGAAATTT TCACGAGTAG TAACCTTTTG CAAGAAGCCC CGCTTTTGAG |
| 1201 | TCAAGGATCT TACCGCTGTT GAGATCCAGT TCGATGTAAC CCACTCGCGC | AGTTCCTAGA ATGGCGACAA CTCTAGGTCA AGCTACATTG GGTGAGCGCG |
| 1251 | ACCCAACCTGA TCCTCAGCAT CTTTACTTT CACCAGCGTT TCTGGGTGAG | TGGGTTGACT AGGAGTCGTA GAAATGAAA GTGGTCGCAA AGACCCACTC  |
| 1301 | CAAAACACAG AAGGCAAAAT GCCGCAAAA AGGGAATAAG GCGACACGG   | GTTTTTGTC TCCCGTTTTA CGCGTTTTT TCCCTTATTC CCGCTGTGCC   |
| 1351 | AAATGTTGAA TACTCATACT CTTCCCTTTT CAATATTATT GAAGCATTTA |                                                        |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |             |            |            |             |            |
|------|-------------|------------|------------|-------------|------------|
|      | TTTACAACCTT | ATGAGTATGA | GAAGGAAAAA | GTTATAATAA  | CTTCGTAAAT |
|      |             |            |            | BsrGI       |            |
|      |             |            |            | ~~~~~       |            |
| 1401 | TCAGGGTTAT  | TGCTCATGA  | GCGGATACAT | ATTTGAATGT  | ACATGAAATT |
|      | AGTCCCAATA  | ACAGAGTACT | CGCCTATGTA | TAAACTTACA  | TGTACTTTAA |
| 1451 | GTAACGTTA   | ATATTTTGTT | AAAATTCGG  | TTAAATTTT   | GTTAAATCAG |
|      | CATTTGCAAT  | TATAAAACAA | TTTTAAGCGC | AATTTAAAAA  | CAATTTAGTC |
| 1501 | CTCATTTTTT  | AACCAATAGG | CCGAAATCGG | CAAAATCCCT  | TATAAATCAA |
|      | GAGTAAAAAA  | TTGGTTATCC | GGCTTTAGCC | GTTTTTAGGA  | ATATTTAGTT |
| 1551 | AAGAATAGAC  | CGAGATAGGG | TTGAGTGTG  | TTCCAGTTTG  | GAACAAGAGT |
|      | TTCTTATCTG  | GCTCTATCCC | AACTCACAA  | AAGTCAAAC   | CTTGTTCTCA |
| 1601 | CCACTATTAA  | AGAACGTGGA | CTCCAACGTC | AAAGGGCGAA  | AAACCGTCTA |
|      | GGTGATAATT  | TCTTGCACCT | GAGGTGCGAG | TTTCCCCGCTT | TTTGGCAGAT |
| 1651 | TCAGGGCGAT  | GGCCCACTAC | GAGAACCATC | ACCCTAATCA  | AGTTTTTTGG |
|      | AGTCCCCGCTA | CCGGGTGATG | CTCTTGGTAG | TGGGATTAGT  | TCAAAAAAAC |

BanII  
~~~~~

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |                                                                                                                  |  |
|------|------------------------------------------------------------------------------------------------------------------|--|
| 1701 | GGTCGAGGTG CCGTAAAGCA CTAAATCGGA ACCCTAAAGG GAGCCCCCGA<br>CCAGCTCCAC GGCATTTTCGT GATTAGCCT TGGGATTTC CTCGGGGGCT  |  |
| 1751 | TTTAGAGCTT GACGGGGAAA GCCGGCGAAC GTGGCGAGAA AGGAAGGGAA<br>AAATCTCGAA CTGCCCCCTT CGGCCGCTTG CACCGCTCTT TCCTTCCCTT |  |
| 1801 | GAAAGCGAAA GGAGCGGGCG CTAGGGCGCT GGCAAGTGTA GCGTCAACG<br>CTTTCGCTTT CCTCGCCCCG GATCCCCGGA CCGTTCACAT CGCCAGTGCG  |  |
| 1851 | TGCGCGTAAC CACCACACC GCCGCGCTTA ATGCGCCGCT ACAGGGCGCG<br>ACGCGCATG GTGGTGTTGG CGGCGCGAAT TACGGCGCGA TGTCCCCGCG   |  |
|      | NheI                                                                                                             |  |
|      | ~~~~~                                                                                                            |  |
| 1901 | TGCTAGCGGA GTGTATACTG GCTTACTATG TTGGCACTGA TGAGGGTGTC<br>ACGATCGCCT CACATATGAC CGAATGATAC AACCGTGACT ACTCCCACAG |  |
|      | XmnI                                                                                                             |  |
|      | ~~~~~                                                                                                            |  |
| 1951 | AGTGAAGTGC TTCATGTGGC AGGAGAAAAA AGGCTGCACC GGTCGTCAG<br>TCACTTCACG AAGTACACCG TCCTCTTTTT TCCGACGTGG CCACGCAGTC  |  |
|      | AgeI                                                                                                             |  |
|      | ~~~~~                                                                                                            |  |
| 2001 | CAGAAATATG GATACAGGAT ATATTCCGCT TCCTCGCTCA CTGACTCGCT<br>GTCTTATACA CTATGTCCTA TATAAGCGA AGGAGCGAGT GACTGAGCGA  |  |



Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|      |                                                                                                                   |
|------|-------------------------------------------------------------------------------------------------------------------|
| 2051 | ACGCTCGGTC GTTCGACTGC GCGAGCGGA AATGGCTTAC GAACGGGGCG<br>TGGGAGCCAG CAAGCTGACG CCGCTCGCCT TTACCGAATG CTGCCCCCGC   |
| 2101 | GAGATTTCCT GGAAGATGCC AGAAGATAC TTAACAGGGA AGTGAGAGGG<br>CTCTAAAGGA CCTTCTACGG TCCTTCTATG AATTGTCCCT TCACTCTCCC   |
| 2151 | CCGCGGCAAA GCCGTTTTC CATAGGCTCC GCCCCCCCTGA CAAGCATCAC<br>GGCGCCCGTT CGGCAAAAAG GTATCCGAGG CGGGGGACT GTTCGTAAGT   |
| 2201 | GAAATCTGAC GCTCAAATCA GTGTGGCGA AACCCGACAG GACTATAAAG<br>CTTTAGACTG CGAGTTTAGT CACCACCGCT TTGGGCTGTC CTGATATTTC   |
| 2251 | ATACCAGGCG TTTCCCCCTG GCGCTCCCT CCTGCGCTCT CCTGTTCCTG<br>TATGCTCCGC AAAGGGGAC CGCCGAGGGA GGACGCGAGA GGACAAGGAC    |
|      | AgeI<br>~~~~~                                                                                                     |
| 2301 | CCTTTCGGTT TACCGGTGTC ATTCCGCTGT TATGGCCGCG TTTGTCTCAT<br>GGAAAGCCAA ATGGCCACAG TAAGGCGACA ATACCGGCGC AACAGAGTA   |
| 2351 | TCCACGCCCTG ACACTCAGTT CCGGGTAGGC AGTTCGCTCC AAGCTGGACT<br>AGGTGCGGAC TGTGAGTCAA GGCCCATCCG TCAAGCGAGG TTCGACCTGA |

Figure 35a: Functional maps and sequences of additional pCAL vector modules and pCAL vectors (continued)

|       |            |             |             |            |             |
|-------|------------|-------------|-------------|------------|-------------|
| 2401  | GTATGCACGA | ACCCCCCGTT  | CAGTCCGACC  | GCTGCGCCTT | ATCCGGTAAC  |
|       | CATACGTGCT | TGGGGGGCAA  | GTCAGGCTGG  | CGACGCGGAA | TAGGCCATTG  |
| 2451  | TATCGTCTTG | AGTCCAACCC  | GGAAAGACAT  | GCAAAAGCAC | CACTGGCAGC  |
|       | ATAGCAGAAC | TCAGGTTGGG  | CCTTCTCTGA  | CGTTTTCGTG | GTGACCGTCG  |
| 2501  | AGCCACTGGT | AATTGATTTA  | GAGGAGTTAG  | TCTTGAAGTC | ATGCCGCCGGT |
|       | TCGGTGACCA | TTAACTAAAT  | CTCCTCAATC  | AGAACTTCAG | TACGCCGGCCA |
| 2551  | TAAGGCTAAA | CTGAAAGGAC  | AAGTTTTAGT  | GACTGCGCTC | CTCCAAGCCA  |
|       | ATTCCGATTT | GACTTTCTCTG | TTCAAAATCA  | CTGACGCGAG | GAGGTTCCGT  |
| 2601  | GTTACCTCGG | TTCAAAGAGT  | TGGTAGCTCA  | GAGAACCTAC | GAAAAACCGC  |
|       | CAATGGAGCC | AAGTTTCTCA  | ACCATCGAGT  | CTCTTGGATG | CTTTTGGCG   |
| 2651  | CCTGCAAGGC | GGTTTTTTTCG | TTTTTCAGAGC | AAGAGATTAC | GCGCAGACCA  |
|       | GGACGTTCCG | CCAAAAAAGC  | AAAAGTCTCG  | TTCTCTAATG | CGCGTCTGGT  |
| BglII |            |             |             |            |             |
| 2701  | AAACGATCTC | AAGAAGATCA  | TCTTATTA    |            |             |
|       | TTTGCTAGAG | TTCTTCTAGT  | AGAAATAT    |            |             |

Figure 35b: List of oligonucleotides used for synthesis of modules

M1: PCR using template

NoVspAatII: TAGACGTC

M2: synthesis

BloxA-A: TATGAGATCTCATAACTTCGTATAATGTACGCTATACG-  
AAGTTAT

BloxA-B: TAATAACTTCGTATAGCATAATTATACGAAGTTATG-  
AGATCTCA

M3: PCR, NoVspAatII as second oligo

XloxS-muta: CATTTTTGCCCTCGTTATCTACGCATGCGATAACTTCGTA-  
TAGCGTACATTATACGAAGTTATTCTAGACATGGTCATAGCTGTTTCCTG

M7-I: PCR

gIIINEW-fow: GGGGGGAATTCGGTGGTGGTGGATCTGCGTGCGCTG-  
AAACGGTTGAAAGTTG

gIIINEW-rev: CCCCCCAAGCTTATCAAGACTCCTTATTACG

M7-II: PCR

gIIIss-fow: GGGGGGGGAATTCGGAGGCGGTTCCGGTGGTGGC

M7-III: PCR

gIIIsupernew-fow: GGGGGGGGAATTCGAGCAGAAGCTGATCTCT-  
GAGGAGGATCTGTAGGGTGGTGGCTCTGGTCCGGTGATTTG

Figure 35b: List of oligonucleotides used for synthesis of modules (continued)

M8: synthesis

lox514-A: CCATAACTTCGTATAATGTACGCTATACGAAGTTATA

lox514-B: AGCTTATAACTTCGTATAGCGTACATTATACGAAGT-  
TATGGCATG

M9II: synthesis

M9II-fow: AGCTTGACCTGTGAAGTGAAAAATGGCGCAGATT-  
GTGCGACATTTTTTTTGTCTGCCGTTTAATTAAAGGGGGGGT

M9II-rev: GTACACCCCCCCCCAGGCCGGCCCCCCCCCCCCCTTTAA-  
TTAAACGGCAGACAAAAAAAATGTCGCACAATCTGCG

M10II: assembly PCR with template

bla-fow: GGGGGGGTGTACATTCAAATATGTATCCGCTCATG

bla-seq4: GGGTTACATCGAACTGGATCTC

bla1-muta: CCAGTTCGATGTAACCCACTCGCGCACCCAACTGATC-  
CTCAGCATCTTTTACTTTCACC

blall-muta: ACTCTAGCTTCCCGGCAACAGTTAATAGACTGGATG-  
GAGGCGG

bla-NEW: CTGTTGCCGGGAAGCTAGAGTAAG

bla-rev: CCCCCCTTAATTAAGGGGGGGGGCCGGCCATTATCAAA-  
AAGGATCTCAAGAAGATCC

M11II/III: PCR, site-directed mutagenesis

Figure 35b: List of oligonucleotides used for synthesis of modules (continued)

f1-fow: GGGGGGGGCTAGCACGCGCCCTGTAGCGGCGCATTA

f1-rev: CCCCCCTGTACATGAAATTGTAAACGTTAATATTTG

f1-t133.muta: GGGCGATGGCCCACTACGAGAACCATCACCTAATC

M12: assembly PCR using template

p15-fow: GGGGGGAGATCTAATAAGATGATCTTCTTGAG

p15-NEWI: GAGTTGGTAGCTCAGAGAACCTACGAAAAACCGCCCTG-  
CAAGGCG

p15-NEWII: GTAGGTTCTCTGAGCTACCAACTC

p15-NEWIII: GTTCCCCCTGGCGGCTCCCTCCTGCGCTCTCCTGTTCT-  
GCC

p15-NEWIV: AGGAGGGAGCCGCCAGGGGGGAAAC

p15-rev: GACATCAGCGCTAGCGGAGTGTATAC

M13: synthesis

BloxXB-A: GATCTCATAACTTCGTATAATGTATGCTATACGAAGTTA-  
TTCA

BloxXB-B: GATCTGAATAACTTCGTATAGCATACATTATACGAAGTTA-  
TGAGA

M14-Ext2: PCR, site-directed mutagenesis

ColEXT2-fow: GGGGGGGGAGATCTGACCAAATCCCTTAACGTGAG

Col-mutal: GGTATCTGCGCTCTGCTGTAGCCAGTTACCTTCGG

Figure 35b: List of oligonucleotides used for synthesis of modules (continued)

Col-rev: CCCCCCGCTAGCCATGTGAGCAAAAGGCCAGCAA

M17: assembly PCR using template

CAT-1: GGGACGTCGGGTGAGGTTCCAAC

CAT-2: CCATACGGAACTCCGGGTGAGCATTTCATC

CAT-3: CCGGAGTTCCGTATGG

CAT-4: ACGTTTAAATCAAACTGG

CAT-5: CCAGTTTTGATTTAAACGTAGCCAATATGGACAACCTTCTTC-

GCCCCCGTTTTCACTATGGGCAAATATT

CAT-6: GGAAGATCTAGCACCAGGCGTTTAAG

M41: assembly PCR using template

LAC1: GAGGCCGGCCATCGAATGGCGCAAAAC

LAC2: CGCGTACCGTCCTCATGGGAGAAAATAATAC

LAC3: CCATGAGGACGGTACGCGACTGGGCGTGGAGCATCTGGTCGCA-

TTGGGTCACCAGCAAATCCGCTGTTAGCTGGCCCATTAAG

LAC4: GTCAGCGGCGGGATATAACATGAGCTGTCCTCGGTATCGTCG

LAC5: GTTATATCCCGCCGCTGACCACCATCAAAC

LAC6: CATCAGTGAATCGGCCAACGCGCGGGGAGAGGCGGTTTGCGT4TTG-

GGAGCCAGGGTGGTTTTTC

LAC7: GGTTAATTAACCTCACTGCCCCGCTTTCAGTCGGGAAACCTGTCGTGCC-

AGCTGCATCAGTGAATCGGCCAAC

M41-MCS-fow: CTAGACTAGTGTTTAAACCGGACCGGGGGGGGGGCTT-

AAGGGGGGGGGGGG

Figure 35b: List of oligonucleotides used for synthesis of modules (continued)

M41-MCS-rev: CTAGCCCCCCCCCCCCCTTAAGCCCCCCCCCGGTCCGGT-TTAAACACTAGT

M41-fow: CTAGACTAGTGTTTAAACCGGACCGGGGGGGGGCTTAA-GGGGGGGGGGGG

M41-rev: CCCCCCTTAAGTGGGCTGCAAAACAAAACGGCCTCC-TGTCAGGAAGCCGCTTTTATCGGGTAGCCTCACTGCCCCGCTTCC

M41-A2: GTTGTGTGCCACGCGGTTAGGAATGTAATTCAGCTCCGC

M41-B1: AACCGCGTGGCACAACAAC

M41-B2: CTCGTTCTACCATCGACACGACCACGCTGGCACCCAGTTG

M41-C1: GTGTCGATGGTAGAACGAAG

M41-CII: CCACAGCAATAGCATCCTGGTCATCCAGCGGATAGTT-AATAATCAGCCCCTGACACGTTGCGCGAG

M41-DI: GACCAGGATGCTATTGCTGTGG

M41-DII: CAGCGCGATTGCTGGTGGCCCAATGCGACCAGATGC

M41-EI: CACCAGCAAATCGCGCTG

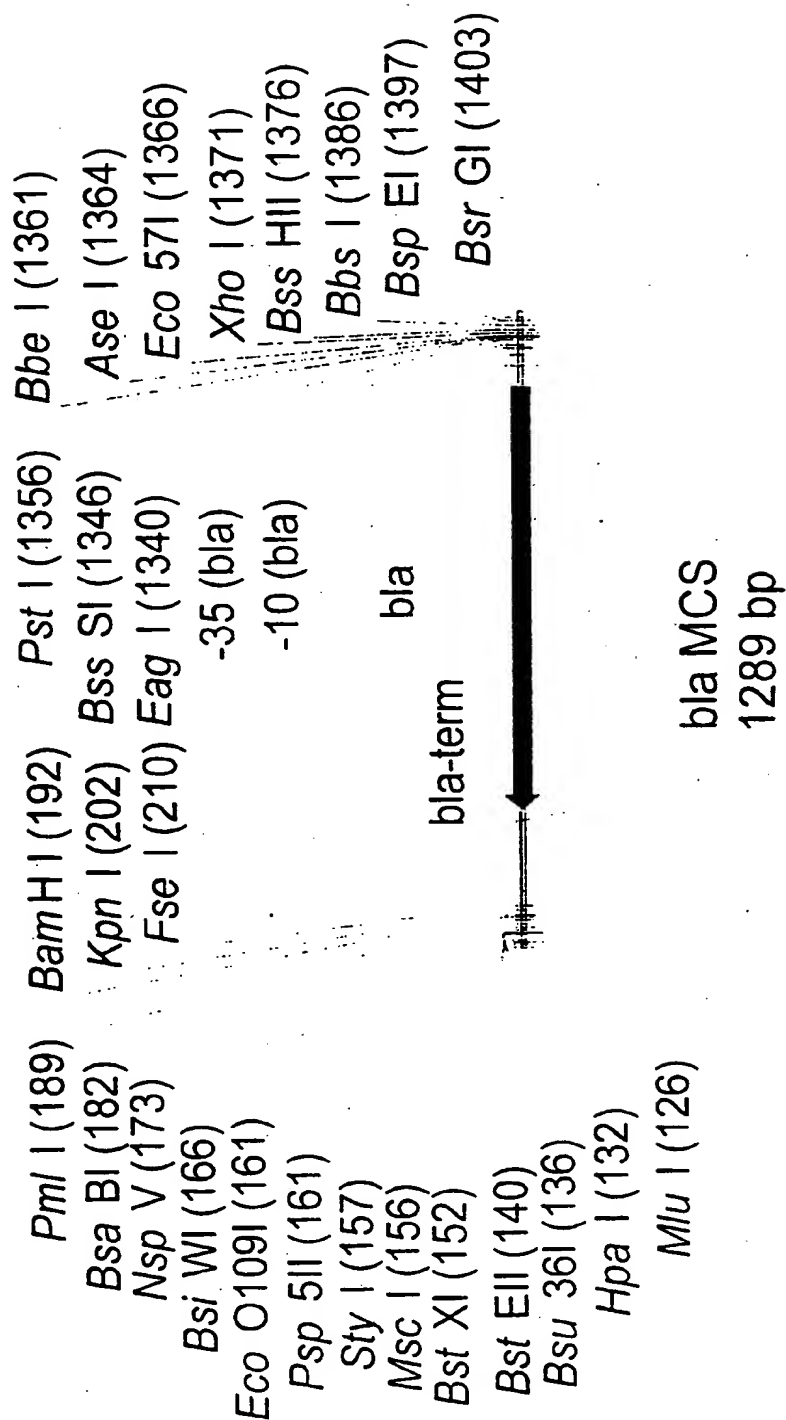
M41-EII: CCCGGACTCGGTAATGGCACGCATTGCGCCCAGCGCC

M41-FI: GCCATTACCGAGTCCGGG

M42: synthesis

Eco-H5-Hind-fow: AATTCCACCATCATCACCATTGACGTCTA

Eco-H5-Hind-rev: AGCTTAGACGTCAATGGTGATGATGGTGG

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module



[illegible]

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

|     |             |            |            |             |             |
|-----|-------------|------------|------------|-------------|-------------|
| 326 | TCACCTAGAT  | CCTTTTAAAT | TAAAAATGAA | GTTTAAATC   | AATCTAAAGT  |
|     | AGTGGATCTA  | GGAAAAATTA | ATTTTACTT  | CAAAATTTAG  | TTAGATTTC   |
| 376 | ATATATGAGT  | AAACTTGGTC | TGACAGTTAC | CAATGCTTAA  | TCAGTGAGGC  |
|     | TATATACTCA  | TTTGAACCAG | ACTGTCAATG | GTTACGAATT  | AGTCACTCCG  |
| 426 | ACCTATCTCA  | GCGATCTGTC | TATTTCTGTC | ATCCATAGTT  | GCCTGACTCC  |
|     | TGGATAGAGT  | CGCTAGACAG | ATAAGCAAG  | TAGGTATCAA  | CGGACTGAGG  |
| 476 | CCGTCGTGTA  | GATAACTACG | ATACGGGAGG | GCTTACCATC  | TGGCCCCAGT  |
|     | GGCAGCACAT  | CTATTGATGC | TATGCCCTCC | CGAATGCTAG  | ACCGGGTCA   |
| 526 | GCTGCAATGA  | TACCGCGAGA | CCCACGCTCA | CCGGCTCCAG  | ATTTATCAGC  |
|     | CGACGTTACT  | ATGGCGCTCT | GGGTGCGAGT | GGCCGAGGTC  | TAAATAGTCG  |
| 576 | AATAAACCAG  | CCAGCCGGAA | GGGCCGAGCG | CAGAAAGTGGT | CCTGCAACTT  |
|     | TTATTTGGTC  | GGTCGGCCTT | CCCGGCTCGC | GTCTTCACCA  | GGACGTTGAA  |
| 626 | TATCCGCCCTC | CATCCAGTCT | ATTAAGTGT  | GCCGGGAAGC  | TAGAGTAAGT  |
|     | ATAGGCGGAG  | GTAGGTCAGA | TAATTGACAA | CGGCCCTTCG  | ATCTCATTTCA |
| 676 | AGTTCGCCCAG | TTAATAGTTT | GCGCAACGTT | GTTGCCCATG  | CTACAGGCAT  |
|     | TCAAGCGGTC  | AATTATCAAA | CGCGTTGCCA | CAACGGTAAC  | GATGTCCGTA  |

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

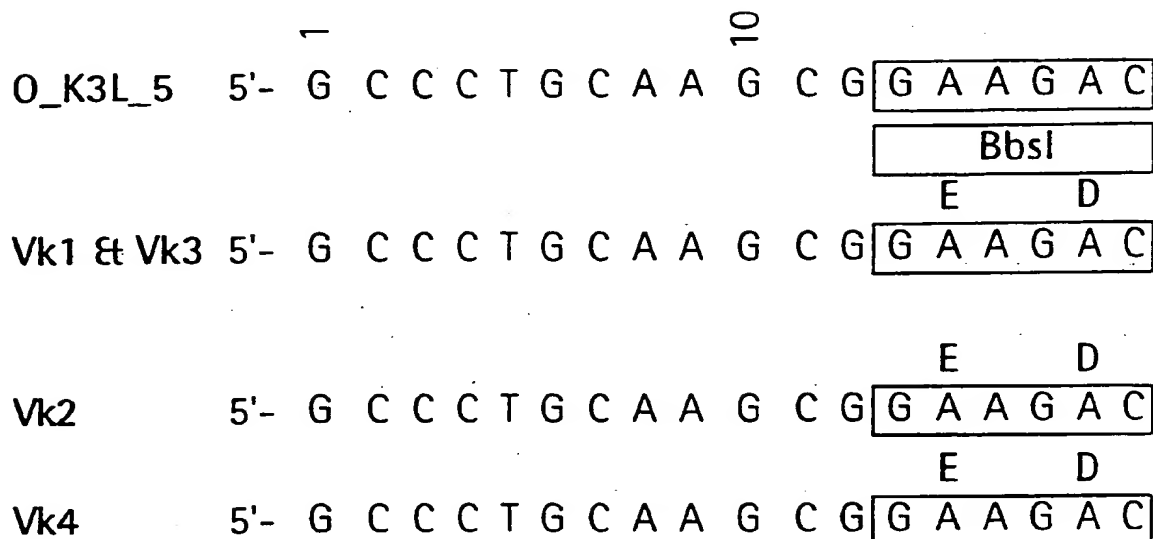
|      |             |            |            |             |             |
|------|-------------|------------|------------|-------------|-------------|
| 726  | CGTGGTGTC   | CGCTCGTCGT | TTGGTATGGC | TTCAATTCAGC | TCCGGTTCCC  |
|      | GCACCACAGT  | GCGAGCAGCA | AACCATACCG | AAGTAAGTCG  | AGGCCAAGGG  |
| 776  | AACGATCAAG  | GCGAGTTACA | TGATCCCCCA | TGTTGTGCAA  | AAAAGCGGTT  |
|      | TTGCTAGTTC  | CGCTCAATGT | ACTAGGGGGT | ACAACACGTT  | TTTTTCGCCAA |
| 826  | AGCTCCCTTCG | GTCCTCCGAT | CGTTGTCAGA | AGTAAGTTGG  | CCGCAGTGTT  |
|      | TCGAGGAAGC  | CAGGAGGCTA | GCAACAGTCT | TCATTCACAC  | GGCGTCACAA  |
| 876  | ATCACTCATG  | GTTATGGCAG | CACTGCATAA | TTCTCTTACT  | GTCATGCCCAT |
|      | TAGTGAGTAC  | CAATACCGTC | GTGACGTATT | AAGAGAATGA  | CAGTACGGTA  |
| 926  | CCGTAAGATG  | CTTTTCTGTG | ACTGGTGAGT | ACTCAACCAA  | GTCATTCTGA  |
|      | GGCATTCTAC  | GAAAGACAC  | TGACCACTCA | TGAGTTGGTT  | CAGTAAGACT  |
| 976  | GAATAGTGTA  | TGCGGCGACC | GAGTTGCTCT | TGCCCGGCCGT | CAATACGGGA  |
|      | CTTATCACAT  | ACGCCGCTGG | CTCAACGAGA | ACGGCCCGCA  | GTTATGCCCT  |
| 1026 | TAATACCGCG  | CCACATAGCA | GAACCTTAAA | AGTGCTCATC  | ATTGGAAAC   |
|      | ATTATGGCGC  | GGTGTAFCGT | CTTGAAATTT | TCACGAGTAG  | TAACCTTTTG  |
| 1076 | GTTCTTCGGG  | GCGAAAACTC | TCAAGGATCT | TACCGCTGTT  | GAGATCCAGT  |
|      | CAAGAAGCCC  | CGCTTTTGAG | AGTTCCTAGA | ATGGCGACAA  | CTCTAGGTCA  |

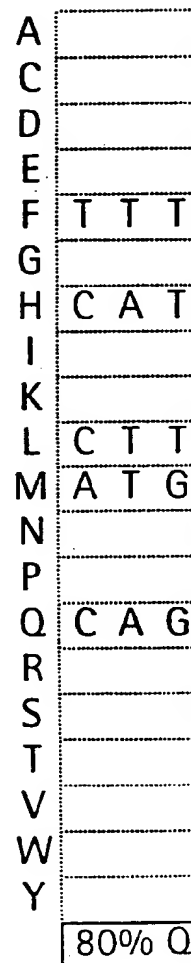
Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

[illegible]

Figure 36: functional map and sequence of  $\beta$ -lactamase-MCS module (continued)

|      |            |             |            |           |
|------|------------|-------------|------------|-----------|
| 1376 | CGCGCTTCAG | CGCTTTGTCT  | TCCGGATGTA | CATGAAATT |
|      | GCGCGAAGTC | GCGAACACAGA | AGCCTACAT  | GTACTTTAA |
|      | Eco57I     | BbsI        |            |           |
|      | ~~~~~      | ~~~~~       |            |           |

Figure 37: Oligo and primer design for V<sub>κ</sub> CDR3 libraries

[illegible]

50

60

# G

A C C T

**G**

A C C T

**G**

A C C T

|       |       |       |       |       |       |
|-------|-------|-------|-------|-------|-------|
| G C T |       |       | G C T |       | G C T |
|       |       |       |       |       |       |
| G A T | G A T | G A T | G A T |       | G A T |
| G A G |       |       | G A G |       | G A G |
| T T T |       |       | T T T |       | T T T |
| G G T | G G T | G G T | G G T |       | G G T |
| C A T |       |       | C A T |       | C A T |
| A . T |       |       | A T T |       | A T T |
| A A G |       |       | A A G |       | A A G |
| C T T |       |       | C T T |       | C T T |
| A T G |       |       | A T G |       | A T G |
| A A T | A A T | A A T | A A T |       | A A T |
|       |       |       | C C T | C C T | C C T |
| C A G |       |       | C A G |       | C A G |
| C G T |       |       | C G T |       | C G T |
| T C T | T C T | T C T | T C T | T C T | T C T |
| A C T |       |       | A C T |       | A C T |
| G T T |       |       | G T T |       | G T T |
| T G G |       |       | T G G |       | T G G |
| T A T | T A T |       | T A T |       | T A T |
| 50% Y |       |       |       | 80% P |       |

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Figure 37: Oligo and primer design for Vκ CDR3 libraries

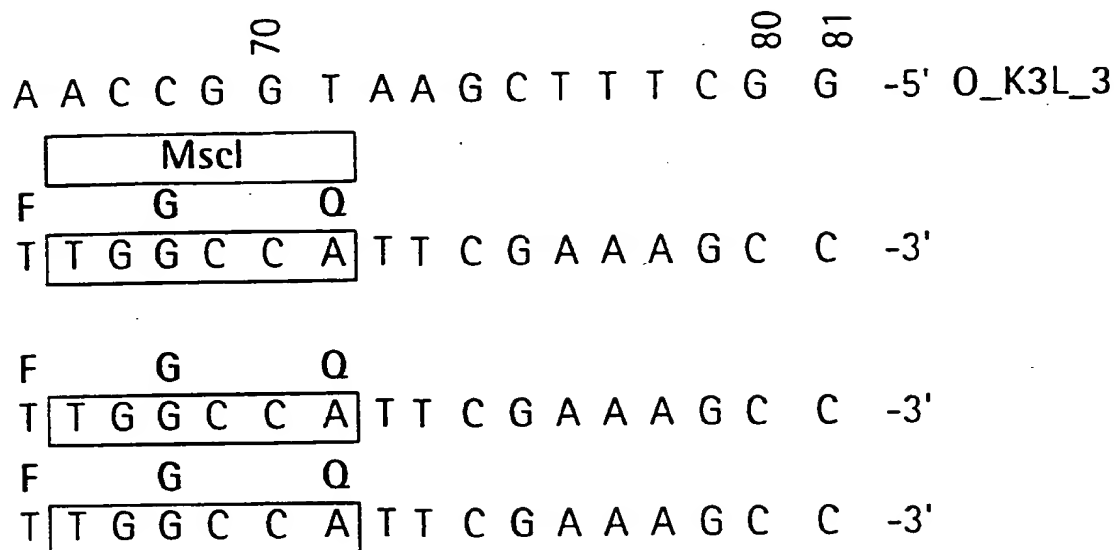
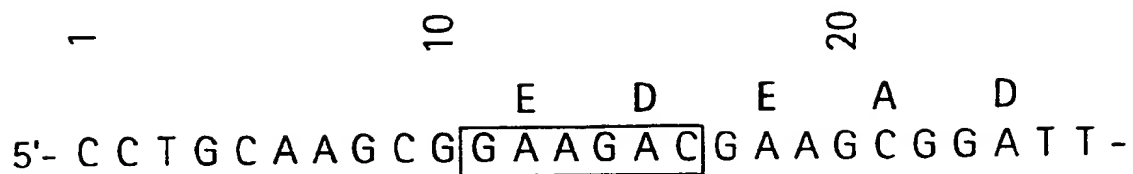


Figure 38: Oligo and primer design for V $\lambda$  CDR3 libraries



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**Figure 38: Oligo and primer design for V $\lambda$  CDR3 libraries**

| 60 |    |    |   | 70  |   |   |   | 80  |    |   |   |
|----|----|----|---|-----|---|---|---|-----|----|---|---|
|    |    |    |   | G   | G | G | T | K   | L  |   |   |
|    |    |    |   | G   | G | C | G | G   | C  | A | C |
| -  | G  | C  | T | gap | G | C | T | gap | G  | C | T |
| G  | A  | T  | G | A   | T | G | A | T   | G  | A | T |
| G  | A  | G  | G | A   | G | G | A | G   | G  | A | G |
| T  | T  | T  | T | T   | T | T | T | T   | T  | T | T |
| G  | G  | T  | G | G   | T | G | G | T   | G  | G | T |
| C  | A  | T  | C | A   | T | C | A | T   | C  | A | T |
| A  | T  | T  | A | T   | T | A | T | T   | A  | T | T |
| A  | A  | G  | A | A   | G | A | A | G   | A  | A | G |
| C  | T  | T  | C | T   | T | C | T | T   | C  | T | T |
| A  | T  | G  | A | T   | G | A | T | G   | A  | T | G |
| A  | A  | T  | A | A   | T | A | A | T   | A  | A | T |
| C  | C  | T  | C | C   | T | C | C | T   | C  | C | T |
| C  | A  | G  | C | A   | G | C | A | G   | C  | A | G |
| C  | G  | T  | C | G   | T | C | G | T   | C  | G | T |
| T  | C  | T  | T | C   | T | T | C | T   | T  | C | T |
| A  | C  | T  | A | C   | T | A | C | T   | A  | C | T |
| G  | T  | T  | G | T   | T | G | T | T   | G  | T | T |
|    |    |    |   |     |   |   |   |     | T  | G | G |
| T  | A  | T  | T | A   | T | T | A | T   | T  | A | T |
| 18 |    |    |   |     |   |   |   |     | 19 |   |   |
| 18 | 18 |    |   |     |   |   |   |     | 19 |   |   |
| 18 | 18 | 18 |   |     |   |   |   |     | 19 |   |   |

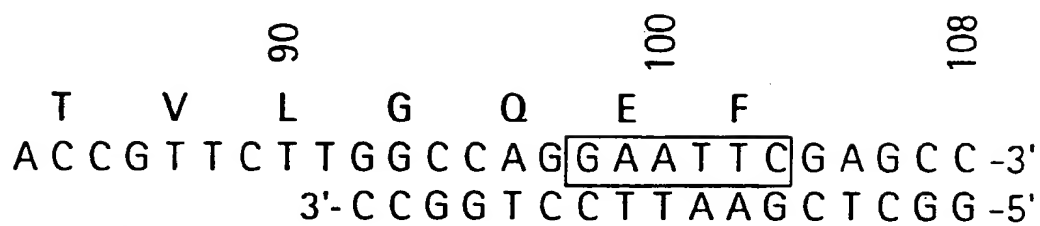
Variability

3.32E+05

5.98E+06

1.08E+08

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Figure 38: Oligo and primer design for V $\lambda$  CDR3 libraries

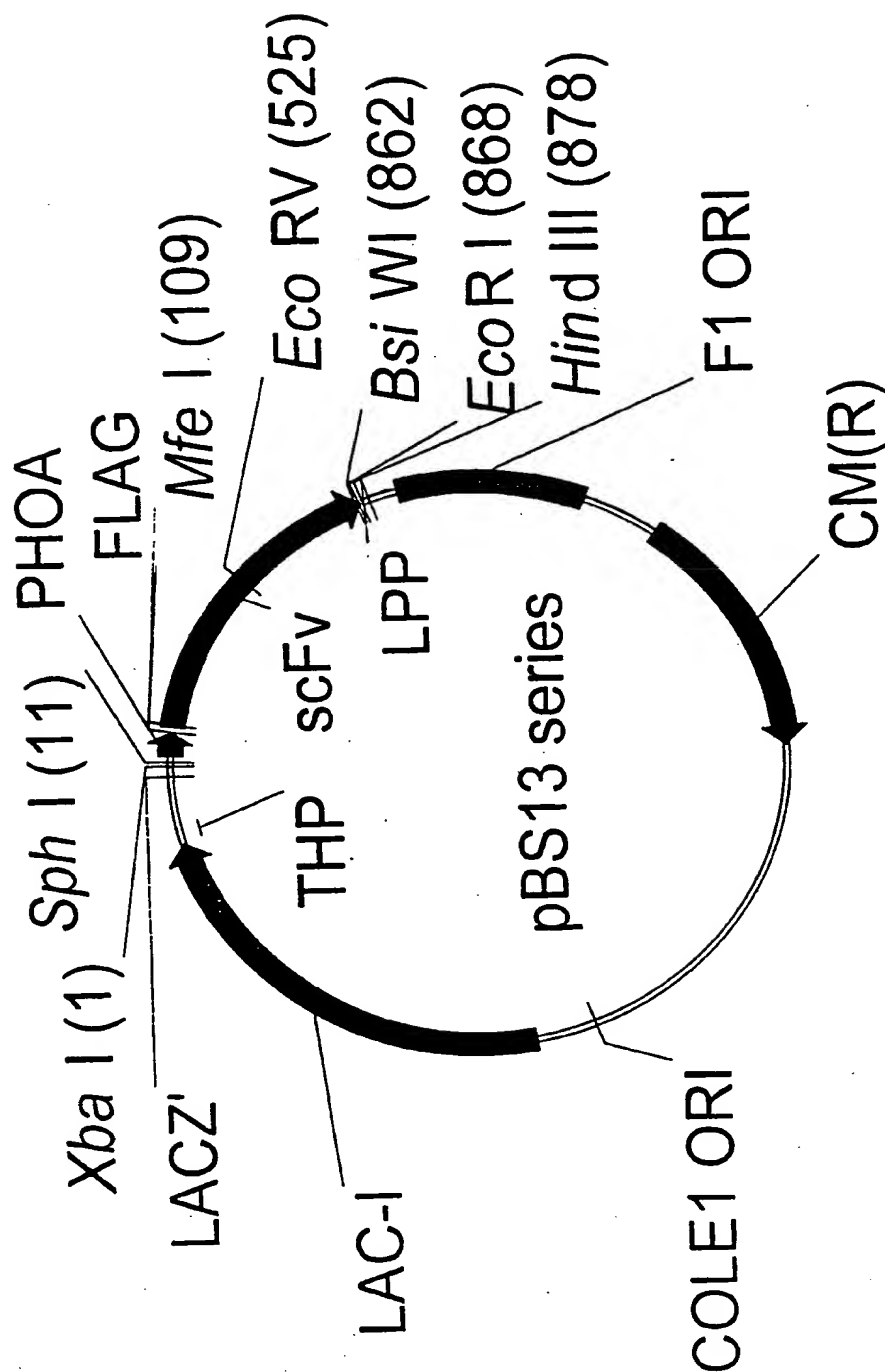


Figure 39: functional map of expression vector series pBS13

Figure 40: Expression data for HuCAL scFvs (pBS13, 30°C)

| % soluble | $\kappa 1$ | $\kappa 2$ | $\kappa 3$ | $\kappa 4$ | $\lambda 1$ | $\lambda 2$ | $\lambda 3$ |
|-----------|------------|------------|------------|------------|-------------|-------------|-------------|
| H1A       | 61%        | 58%        | 52%        | 42%        | 90%         | 61%         | 60%         |
| H1B       | 39%        | 48%        | 66%        | 48%        | 47%         | 39%         | 36%         |
| H2        | 47%        | 57%        | 46%        | 49%        | 37%         | 36%         | 45%         |
| H3        | 85%        | 67%        | 76%        | 61%        | 80%         | 71%         | 83%         |
| H4        | 69%        | 52%        | 51%        | 44%        | 45%         | 33%         | 42%         |
| H5        | 49%        | 49%        | 46%        | 67%        | 54%         | 46%         | 47%         |
| H6        | 90%        | 58%        | 54%        | 47%        | 45%         | 50%         | 51%         |

| Total amount compared to H3 $\kappa 2$ | $\kappa 1$ | $\kappa 2$ | $\kappa 3$ | $\kappa 4$ | $\lambda 1$ | $\lambda 2$ | $\lambda 3$ |
|----------------------------------------|------------|------------|------------|------------|-------------|-------------|-------------|
| H1A                                    | 289%       | 94%        | 166%       | 272%       | 20%         | 150%        | 78%         |
| H1B                                    | 219%       | 122%       | 89%        | 139%       | 117%        | 158%        | 101%        |
| H2                                     | 186%       | 223%       | 208%       | 182%       | 126%        | 60%         | 97%         |
| H3                                     | 50%        |            | 71%        | 54%        | 59%         | 130%        | 47%         |
| H4                                     | 37%        | 55%        | 60%        | 77%        | 195%        | 107%        | 251%        |
| H5                                     | 98%        | 201%       | 167%       | 83%        | 93%         | 128%        | 115%        |
| H6                                     | 65%        | 117%       | 89%        | 109%       | 299%        | 215%        | 278%        |

Figure 40: Expression data for HuCAL scfvs (pBS13, 30°C)

| Soluble amount<br>compared to H3k2 | $\kappa 1$ | $\kappa 2$ | $\kappa 3$ | $\kappa 4$ | $\lambda 1$ | $\lambda 2$ | $\lambda 3$ |
|------------------------------------|------------|------------|------------|------------|-------------|-------------|-------------|
| H1A                                | 191%       | 88%        | 121%       | 122%       | 26%         | 211%        | 76%         |
| H1B                                | 124%       | 95%        | 83%        | 107%       | 79%         | 142%        | 59%         |
| H2                                 | 126%       | 204%       | 139%       | 130%       | 66%         | 50%         | 70%         |
| H3                                 | 63%        | -          | 81%        | 49%        | 69%         | 143%        | 61%         |
| H4                                 | 40%        | 47%        | 49%        | 54%        | 95%         | 55%         | 125%        |
| H5                                 | 69%        | 158%       | 116%       | 80%        | 72%         | 84%         | 84%         |
| H6                                 | 85%        | 122%       | 87%        | 77%        | 162%        | 162%        | 212%        |
|                                    | McPC       |            |            |            |             |             |             |
| soluble                            | 38%        |            |            |            |             |             |             |
| %H3k2 total                        | 117%       |            |            |            |             |             |             |
| %H3k2 soluble                      | 69%        |            |            |            |             |             |             |